

In the Specification:

Please amend the specification as shown.

Please amend the paragraph on page 18, lines 6-27, as follows:

In another embodiment of the invention, said nucleotides are linked to each other by means of a phosphorothioate group, such as all nucleotides being linked to each other by means of a phosphorothioate group. An interesting embodiment of the invention is directed to compounds of SEQ NO 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120, 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141, 142, 143, and 144 wherein each linkage group within each compound is a phosphorothioate group. Such modifications is denoted by the subscript S. Alternatively stated, one aspect of the invention is directed to compounds of SEQ NO ~~2_{A5}, 3_{A5}, 4_{A5}, 5_{A5}, 6_{S5}, 7_{S5}, 8_{S5}, 9_{A5}, 10_{A5}, 11_{A5}, 12_{A5}, 13_{A5}, 14_{A5}, 15_{A5}, 16_{A5}, 17_{A5}, 18_{A5}, 19_{A5}, 20_{A5}, 21_{A5}, 22_{A5}, 23_{A5}, 24_{A5}, 25_{A5}, 26_{A5}, 27_{A5}, 28_{A5}, 29_{A5}, 30_{A5}, 31_{A5}, 32_{A5}, 33_{A5}, 34_{A5}, 35_{A5}, 36_{A5}, 37_{S5}, 38_{A5}, 39_{A5}, 40_{A5}, 41_{A5}, 42_{A5}, 43_{A5}, 44_{A5}, 45_{A5}, 46_{A5}, 47_{A5}, 48_{A5}, 49_{A5}, 50_{A5}, 51_{A5}, 52_{A5}, 53_{A5}, 54_{A5}, 55_{A5}, 56_{A5}, 57_{A5}, 58_{A5}, 59_{A5}, 60_{A5}, 61_{A5}, 62_{A5}, 63_{A5}, 64_{A5}, 65_{A5}, 66_{A5}, 67_{A5}, 68_{A5}, 69_{A5}, 70_{A5}, 71_{A5}, 72_{A5}, 73_{A5}, 74_{A5}, 75_{A5}, 76_{A5}, 77_{A5}, 78_{A5}, 79_{A5}, 80_{A5}, 81_{A5}, 82_{A5}, 83_{A5}, 84_{A5}, 85_{A5}, 86_{A5}, 87_{A5}, 88_{A5}, 89_{A5}, 90_{A5}, 91_{A5}, 92_{A5}, 93_{A5}, 94_{A5}, 95_{A5}, 96_{A5}, 97_{A5}, 98_{A5}, 99_{A5}, 100_{A5}, 101_{A5}, 102_{A5}, 103_{A5}, 104_{A5}, 105_{A5}, 106_{A5}, 107_{A5}, 108_{A5}, 109_{A5}, 101_{A5}, 102_{A5}, 103_{A5}, 104_{A5}, 105_{A5}, 106_{A5}, 107_{A5}, 108_{A5}, 109_{A5}, 110_{A5}, 111_{A5}, 112_{A5}, 113_{A5}, 114_{A5}, 115_{A5}, 116_{A5}, 117_{A5}, 118_{A5}, 119_{A5}, 120_{A5}, 121_{A5}, 122_{A5}, 123_{A5}, 124_{A5}, 125_{A5}, 126_{A5}, 127_{A5}, 128_{A5}, 129_{A5}, 130_{A5}, 131_{A5}, 132_{A5}, 133_{A5}, 134_{A5}, 135_{A5}, 136_{A5}, 137_{A5}, 138_{A5}, 139_{A5}, 140_{A5}, 141_{A5}, 142_{A5}, 143_{A5} and 144_{A5}, 147, 151, 155, 159, 163, 167, 171, 175, 179, 183, 187, 191, 195, 199, 204, 208, 212, 216, 220, 224, 228, 232, 236, 240, 244, 248, 252, 256, 260, 264, 268, 272, 276, 280, 284, 288, 292, 296, 300, 304, 308, 312, 316, 320, 324, 328, 332, 336, 340, 344, 348, 352,~~

356, 360, 364, 368, 372, 376, 380, 384, 388, 392, 396, 400, 404, 408, 412, 416, 420, 424, 428, 432, 436, 440, 444, 448, 452, 456, 460, 464, 468, 472, 476, 480, 484, 488, 492, 496, 500, 504, 508, 512, 516, 520, 524, 528, 532, 536, 540, 544, 548, 552, 556, 560, 564, 568, 572, 576, 580, 584, 588, 592, 596, 600, 604, 608, 612, 616, 620, 624, 628, 632, 636, 640, 644, 648, 652, 656, 660, 664, 668, 672, 676, 680, 684, 688, 692, 696, 700, 704, 708, 712 and 716.

Please amend the paragraph on page 18, lines 29-30, as follows:

A preferred subset of embodiments of the invention are compounds comprising sequences of the formula ~~2_A, 4_A, 6_A, 9_A, 15_A, 118_A, 120_A, 123_A, 128_A, 129_A, and 131_A~~ SEQ ID NOS 147, 155, 163, 175, 199, 612, 620, 632, 652, 656 and 664.

Please amend the paragraph on page 18, line 32, to page 19, line 2, as follows:

A further aspect of the invention is directed to compounds of SEQ NOS ~~2_B, 3_B, 4_B, 5_B, 6_S, 7_S, 8_B, 9_B, 10_B, 11_B, 12_B, 13_B, 14_B, 15_B, 16_B, 17_B, 18_B, 19_B, 20_B, 21_B, 22_B, 23_B, 24_B, 25_B, 26_B, 27_B, 28_B, 29_B, 30_B, 31_B, 32_B, 33_B, 34_B, 35_B, 36_B, 37_S, 38_B, 39_B, 40_B, 41_B, 42_B, 43_B, 44_B, 45_B, 46_B, 47_B, 48_B, 49_B, 50_B, 51_B, 52_B, 53_B, 54_B, 55_B, 56_B, 57_B, 58_B, 59_B, 60_B, 61_B, 62_B, 63_B, 64_B, 65_B, 66_B, 67_B, 68_B, 69_B, 70_B, 71_B, 72_B, 73_B, 74_B, 75_B, 76_B, 77_B, 78_B, 79_B, 80_B, 81_B, 82_B, 83_B, 84_B, 85_B, 86_B, 87_B, 88_B, 89_B, 90_B, 91_B, 92_B, 93_B, 94_B, 95_B, 96_B, 97_B, 98_B, 99_B, 100_B, 101_B, 102_B, 103_B, 104_B, 105_B, 106_B, 107_B, 108_B, 109_B, 110_B, 111_B, 112_B, 113_B, 114_B, 115_B, 116_B, 117_B, 118_B, 119_B, 120_B, 121_B, 122_B, 123_B, 124_B, 125_B, 126_B, 127_B, 128_B, 129_B, 130_B, 131_B, 132_B, 133_B, 134_B, 135_B, 136_B, 137_B, 138_B, 139_B, 140_B, 141_B, 142_B, 143_B, and 144_B~~ 148, 152, 156, 160, 164, 168, 172, 176, 180, 184, 188, 192, 196, 200, 205, 209, 213, 217, 221, 225, 229, 233, 237, 241, 245, 249, 253, 257, 261, 265, 269, 273, 277, 281, 285, 289, 293, 297, 301, 305, 309, 313, 317, 321, 325, 329, 333, 337, 341, 345, 349, 353, 357, 361, 365, 369, 373, 377, 381, 385, 389, 393, 397, 401, 405, 409, 413, 417, 421, 425, 429, 433, 437, 441, 445, 449, 453, 457, 461, 465, 469,

473, 477, 481, 485, 489, 493, 497, 501, 505, 509, 513, 517, 521, 525, 529, 533, 537, 541, 545, 549, 553, 557, 561, 565, 569, 573, 577, 581, 585, 589, 593, 597, 601, 605, 609, 613, 617, 621, 625, 629, 633, 637, 641, 645, 649, 653, 657, 661, 665, 669, 673, 677, 681, 685, 689, 693, 697, 701, 705, 709, 713 and 717.

Please amend the paragraph on page 19, lines 4-5, as follows:

A preferred subset of embodiments of the invention are compounds comprising sequences of the SEQ ID NOS ~~formula 118_B, 119_B, 120_B, 121_B, 122_B, 123_B, 128_B, 129_B, 130_B, and 131_B~~ 613, 617, 621, 625, 629, 633, 653, 657, 661 and 665.

Please amend the paragraph on page 19, lines 7-16, as follows:

A further aspect of the invention is directed to compounds of SEQ NOS ~~2_C, 3_C, 4_C, 5_C, 6_S, 7_S, 8_C, 9_C, 10_C, 11_C, 12_C, 13_C, 14_C, 15_C, 16_C, 17_C, 18_C, 19_C, 20_C, 21_C, 22_C, 23_C, 24_C, 25_C, 26_C, 27_C, 28_C, 29_C, 30_C, 31_C, 32_C, 33_C, 34_C, 35_C, 36_C, 37_S, 38_C, 39_C, 40_C, 41_C, 42_C, 43_C, 44_C, 45_C, 46_C, 47_C, 48_C, 49_C, 50_C, 51_C, 52_C, 53_C, 54_C, 55_C, 56_C, 57_C, 58_C, 59_C, 60_C, 61_C, 62_C, 63_C, 64_C, 65_C, 66_C, 67_C, 68_C, 69_C, 70_C, 71_C, 72_C, 73_C, 74_C, 75_C, 76_C, 77_C, 78_C, 79_C, 80_C, 81_C, 82_C, 83_C, 84_C, 85_C, 86_C, 87_C, 88_C, 89_C, 90_C, 91_C, 92_C, 93_C, 94_C, 95_C, 96_C, 97_C, 98_C, 99_C, 100_C, 101_C, 102_C, 103_C, 104_C, 105_C, 106_C, 107_C, 108_C, 109_C, 101_C, 102_C, 103_C, 104_C, 105_C, 106_C, 107_C, 108_C, 109_C, 110_C, 111_C, 112_C, 113_C, 114_C, 115_C, 116_C, 117_C, 118_C, 119_C, 120_C, 121_C, 122_C, 123_C, 124_C, 125_C, 126_C, 127_C, 128_C, 129_C, 130_C, 131_C, 132_C, 133_C, 134_C, 135_C, 136_C, 137_C, 138_C, 139_C, 140_C, 141_C, 142_C, 143_C, and 144_C~~ 149, 153, 157, 161, 165, 169, 173, 177, 181, 185, 189, 193, 197, 201, 206, 210, 214, 218, 222, 226, 230, 234, 238, 242, 246, 250, 254, 258, 262, 266, 270, 274, 278, 282, 286, 290, 294, 298, 302, 306, 310, 314, 318, 322, 326, 330, 334, 338, 342, 346, 350, 354, 358, 362, 366, 370, 374, 378, 382, 386, 390, 394, 398, 402, 406, 410, 414, 418, 422, 426, 430, 434, 438, 442, 446, 450, 454, 458, 462, 466, 470, 474, 478, 482, 486, 490, 494, 498, 502, 506, 510, 514, 518, 522, 526, 530, 534, 538, 542,

546, 550, 554, 558, 562, 566, 570, 574, 578, 582, 586, 590, 594, 598, 602, 606, 610, 614, 618, 622, 626, 630, 634, 638, 642, 646, 650, 654, 658, 662, 666, 670, 674, 678, 682, 686, 690, 694, 698, 702, 706, 710, 714 and 718.

Please amend the paragraph on page 19, lines 18-27, as follows:

A further aspect of the invention is directed to compounds of SEQ NOS ~~2_D, 3_D, 4_D, 5_D, 6_S, 7_S, 8_D, 9_D, 10_D, 11_D, 12_D, 13_D, 14_D, 15_D, 16_D, 17_D, 18_D, 19_D, 20_D, 21_D, 22_D, 23_D, 24_D, 25_D, 26_D, 27_D, 28_D, 29_D, 30_D, 31_D, 32_D, 33_D, 34_D, 35_D, 36_D, 37_S, 38_D, 39_D, 40_D, 41_D, 42_D, 43_D, 44_D, 45_D, 46_D, 47_D, 48_D, 49_D, 50_D, 51_D, 52_D, 53_D, 54_D, 55_D, 56_D, 57_D, 58_D, 59_D, 60_D, 61_D, 62_D, 63_D, 64_D, 65_D, 66_D, 67_D, 68_D, 69_D, 70_D, 71_D, 72_D, 73_D, 74_D, 75_D, 76_D, 77_D, 78_D, 79_D, 80_D, 81_D, 82_D, 83_D, 84_D, 85_D, 86_D, 87_D, 88_D, 89_D, 90_D, 91_D, 92_D, 93_D, 94_D, 95_D, 96_D, 97_D, 98_D, 99_D, 100_D, 101_D, 102_D, 103_D, 104_D, 105_D, 106_D, 107_D, 108_D, 109_D, 101_D, 102_D, 103_D, 104_D, 105_D, 106_D, 107_D, 108_D, 109_D, 110_D, 111_D, 112_D, 113_D, 114_D, 115_D, 116_D, 117_D, 118_D, 119_D, 120_D, 121_D, 122_D, 123_D, 124_D, 125_D, 126_D, 127_D, 128_D, 129_D, 130_D, 131_D, 132_D, 133_D, 134_D, 135_D, 136_D, 137_D, 138_D, 139_D, 140_D, 141_D, 142_D, 143_D, and 144_D.~~ 150, 154, 158, 162, 166, 170, 174, 178, 182, 186, 190, 194, 198, 202, 207, 211, 215, 219, 223, 227, 231, 235, 239, 243, 247, 251, 255, 259, 263, 267, 271, 275, 279, 283, 287, 291, 295, 299, 303, 307, 311, 315, 319, 323, 327, 331, 335, 339, 343, 347, 351, 355, 359, 363, 367, 371, 375, 379, 383, 387, 391, 395, 399, 403, 407, 411, 415, 419, 423, 427, 431, 435, 439, 443, 447, 451, 455, 459, 463, 467, 471, 475, 479, 483, 487, 491, 495, 499, 503, 507, 511, 515, 519, 523, 527, 531, 535, 539, 543, 547, 551, 555, 559, 563, 567, 571, 575, 579, 583, 587, 591, 595, 599, 603, 607, 611, 615, 619, 623, 627, 631, 635, 639, 643, 647, 651, 655, 659, 663, 667, 671, 675, 679, 683, 687, 691, 695, 699, 703, 707, 711, 715 and 719.

Please amend the paragraph on page 19, lines 29-38, as follows:

A further aspect of the invention is directed to compounds of ~~SEQ NO 2_E, 3_E, 4_E, 5_E, 6_S, 7_S, 8_E, 9_E, 10_E, 11_E, 12_E, 13_E, 14_E, 15_E, (SEQ ID NO: 203) 16_E, 17_E, 18_E, 19_E, 20_E, 21_E, 22_E, 23_E, 24_E, 25_E, 26_E, 27_E, 28_E, 29_E, 30_E, 31_E, 32_E, 33_E, 34_E, 35_E, 36_E, 37_S, 38_E, 39_E, 40_E, 41_E, 42_E, 43_E, 44_E, 45_E, 46_E, 47_E, 48_E, 49_E, 50_E, 51_E, 52_E, 53_E, 54_E, 55_E, 56_E, 57_E, 58_E, 59_E, 60_E, 61_E, 62_E, 63_E, 64_E, 65_E, 66_E, 67_E, 68_E, 69_E, 70_E, 71_E, 72_E, 73_E, 74_E, 75_E, 76_E, 77_E, 78_E, 79_E, 80_E, 81_E, 82_E, 83_E, 84_E, 85_E, 86_E, 87_E, 88_E, 89_E, 90_E, 91_E, 92_E, 93_E, 94_E, 95_E, 96_E, 97_E, 98_E, 99_E, 100_E, 101_E, 102_E, 103_E, 104_E, 105_E, 106_E, 107_E, 108_E, 109_E, 101_E, 102_E, 103_E, 104_E, 105_E, 106_E, 107_E, 108_E, 109_E, 110_E, 111_E, 112_E, 113_E, 114_E, 115_E, 116_E, 117_E, 118_E, 119_E, 120_E, 121_E, 122_E, 123_E, 124_E, 125_E, 126_E, 127_E, 128_E, 129_E, 130_E, 131_E, 132_E, 133_E, 134_E, 135_E, 136_E, 137_E, 138_E, 139_E, 140_E, 141_E, 142_E, 143_E, and 144_E.~~

Please amend the paragraph on page 23 line 26 to page 24, line 12 as follows:

In a suitable embodiment, the subsequence is SEQ ID NO: 147 2a. In a suitable embodiment, the subsequence is SEQ ID NO: 151 3a. In a suitable embodiment, the subsequence is SEQ ID NO: 155 4a. In a suitable embodiment, the subsequence is SEQ ID NO: 159 5a. In a suitable embodiment, the subsequence is SEQ ID NO: 163 6a. In a suitable embodiment, the subsequence is SEQ ID NO: 167 7a. In a suitable embodiment, the subsequence is SEQ ID NO: 171 8a. In a suitable embodiment, the subsequence is SEQ ID NO: 175 9a. In a suitable embodiment, the subsequence is SEQ ID NO: 179 10a. In a suitable embodiment, the subsequence is SEQ ID NO: 183 11a. In a suitable embodiment, the subsequence is SEQ ID NO: 187 12a. In a suitable embodiment, the subsequence is SEQ ID NO: 191 13a. In a suitable embodiment, the subsequence is SEQ ID NO: 195 14a. In a suitable embodiment, the subsequence is SEQ ID NO: 199 15a. In a suitable embodiment, the subsequence is SEQ ID NO: 608 117a. In a suitable embodiment, the subsequence is SEQ ID NO: 612 118a. In a suitable embodiment, the subsequence is SEQ ID NO: 616 119a. In a suitable embodiment, the subsequence is SEQ ID NO: 620 120a. In a suitable embodiment, the subsequence is SEQ

ID NO: 624 121a. In a suitable embodiment, the subsequence is SEQ ID NO: 628 122a. In a suitable embodiment, the subsequence is SEQ ID NO: 632 123a. In a suitable embodiment, the subsequence is SEQ ID NO: 636 124a. In a suitable embodiment, the subsequence is SEQ ID NO: 640 125a. In a suitable embodiment, the subsequence is SEQ ID NO: 644 126a. In a suitable embodiment, the subsequence is SEQ ID NO: 648 127a. In a suitable embodiment, the subsequence is SEQ ID NO: 652 128a. In a suitable embodiment, the subsequence is SEQ ID NO: 656 129a. In a suitable embodiment, the subsequence is SEQ ID NO: 660 130a. In a suitable embodiment, the subsequence is SEQ ID NO: 664 131a. In a suitable embodiment, the subsequence is SEQ ID NO: 668 132a. In a suitable embodiment, the subsequence is SEQ ID NO: 672 133a. In the immediately aforementioned individual suitable embodiments wherein the subsequence is one selected from SEQ ID NOS: 2a-144a-148, 152, 156, 160, 164, 168, 172, 176, 180, 184, 188, 192, 196, 200, 205, 209, 213, 217, 221, 225, 229, 233, 237, 241, 245, 249, 253, 257, 261, 265, 269, 273, 277, 281, 285, 289, 293, 297, 301, 305, 309, 313, 317, 321, 325, 329, 333, 337, 341, 345, 349, 353, 357, 361, 365, 369, 373, 377, 381, 385, 389, 393, 397, 401, 405, 409, 413, 417, 421, 425, 429, 433, 437, 441, 445, 449, 453, 457, 461, 465, 469, 473, 477, 481, 485, 489, 493, 497, 501, 505, 509, 513, 517, 521, 525, 529, 533, 537, 541, 545, 549, 553, 557, 561, 565, 569, 573, 577, 581, 585, 589, 593, 597, 601, 605, 609, 613, 617, 621, 625, 629, 633, 637, 641, 645, 649, 653, 657, 661, 665, 669, 673, 677, 681, 685, 689, 693, 697, 701, 705, 709, 713 and 717, the 3' end LNA of the subsequence may suitably be replaced by the corresponding nucleotide.

Please amend the paragraph on page 24 line 21 to page 25, line 9, as follows:

In a suitable embodiment, the compound consists of SEQ ID NO: 147 2a. In a suitable embodiment, the compound consists of SEQ ID NO: 151 3a. In a suitable embodiment, the compound consists of SEQ ID NO: 155 4a. In a suitable embodiment, the compound consists of SEQ ID NO: 159 5a. In a suitable embodiment, the compound consists of SEQ ID NO: 163 6a. In a suitable embodiment, the compound consists of SEQ ID NO: 167 7a. In a suitable

embodiment, the compound consists of SEQ ID NO: 171 8a. In a suitable embodiment, the compound consists of SEQ ID NO: 175 9a. In a suitable embodiment, the compound consists of SEQ ID NO: 179 10a. In a suitable embodiment, the compound consists of SEQ ID NO: 183 11a. In a suitable embodiment, the compound consists of SEQ ID NO: 187 12a. In a suitable embodiment, the compound consists of SEQ ID NO: 191 13a. In a suitable embodiment, the compound consists of SEQ ID NO: 195 14a. In a suitable embodiment, the compound consists of SEQ ID NO: 199 15a. In a suitable embodiment, the compound consists of SEQ ID NO: 608 117a. In a suitable embodiment, the compound consists of SEQ ID NO: 612 118a. In a suitable embodiment, the compound consists of SEQ ID NO: 616 119a. In a suitable embodiment, the compound consists of SEQ ID NO: 620 120a. In a suitable embodiment, the compound consists of SEQ ID NO: 624 121a. In a suitable embodiment, the compound consists of SEQ ID NO: 628 122a. In a suitable embodiment, the compound consists of SEQ ID NO: 632 123a. In a suitable embodiment, the compound consists of SEQ ID NO: 636 124a. In a suitable embodiment, the compound consists of SEQ ID NO: 640 125a. In a suitable embodiment, the compound consists of SEQ ID NO: 644 126a. In a suitable embodiment, the compound consists of SEQ ID NO: 648 127a. In a suitable embodiment, the compound consists of SEQ ID NO: 652 128a. In a suitable embodiment, the compound consists of SEQ ID NO: 656 129a. In a suitable embodiment, the compound consists of SEQ ID NO: 660 130a. In a suitable embodiment, the compound consists of SEQ ID NO: 664 131a. In a suitable embodiment, the compound consists of SEQ ID NO: 668 132a. In a suitable embodiment, the compound consists of SEQ ID NO: 672 133a. In the immediately aforementioned individual suitable embodiments wherein the compound is one selected from SEQ ID NOS: 2a-144a-147, 151, 155, 159, 163, 167, 171, 175, 179, 183, 187, 191, 195, 199, 204, 208, 212, 216, 220, 224, 228, 232, 236, 240, 244, 248, 252, 256, 260, 264, 268, 272, 276, 280, 284, 288, 292, 296, 300, 304, 308, 312, 316, 320, 324, 328, 332, 336, 340, 344, 348, 352, 356, 360, 364, 368, 372, 376, 380, 384, 388, 392, 396, 400, 404, 408, 412, 416, 420, 424, 428, 432, 436, 440, 444, 448, 452, 456, 460, 464, 468, 472, 476, 480, 484, 488, 492, 496, 500, 504, 508, 512, 516, 520, 524, 528, 532,

536, 540, 544, 548, 552, 556, 560, 564, 568, 572, 576, 580, 584, 588, 592, 596, 600, 604, 608, 612, 616, 620, 624, 628, 632, 636, 640, 644, 648, 652, 656, 660, 664, 668, 672, 676, 680, 684, 688, 692, 696, 700, 704, 708, 712 and 716, the 3' end LNA of the compound may suitably be replaced by the corresponding nucleotide.

Please amend the paragraph on page 63 lines 8-15 as follows:

First strand synthesis was performed using OmniScript Reverse Transcriptase kit (cat# 205113, Qiagen) according to the manufacturers instructions.
For each sample 0.5 µg total RNA was adjusted to 12 µl each with RNase free H₂O and mixed with 2 µl poly (dT)₁₂₋₁₈ (SEQ ID NO: 741) (2.5 µg/ml) (Life Technologies, GibcoBRL, Roskilde, DK), 2 µl dNTP mix (5 mM each dNTP), 2 µl 10x Buffer RT, 1 µl RNAguard™Rnase INHIBITOR (33.3U/ml), (cat# 27-0816-01, Amersham Pharmacia Biotech, Hørsholm, DK) and 1 µl OmniScript Reverse Transcriptase (4 U/µl) followed by incubation at 37°C for 60 minutes and heat inactivation of the enzyme at 93°C for 5 minutes.

Please amend the paragraph on page 64 lines 9-18, as follows:

For human Survivin the PCR primers were:

Assay 1

forward primer: 5' caggtccccgctttcttg 3' (SEQ ID NO: 727) (final concentration in the assay; 0.6 µM)

reverse primer: 5' ggaggaggcgcaatcaaa 3' (SEQ ID NO: 728) (final concentration in the assay; 0.6 µM) and the PCR probe was: 5' FAM- ccatactctacgccagacttcagcc-TAMRA 3' (SEQ ID NO: 729) (final concentration in the assay; 0.1 µM) Assay 2

forward primer: 5' aaggaccaccgcatctctaca 3' (SEQ ID NO: 730) (final concentration in the assay; 0.9 µM)

reverse primer: 5' ccaagtctggctcgttctcagt 3' (SEQ ID NO: 731) (final concentration in the assay; 0.6 μ M) and the PCR probe was: 5' FAM- cgaggctggcttcacccactgcc -TAMRA 3' (SEQ ID NO: 732) (final concentration in the assay; 0.1 μ M)

Please amend the paragraph on page 64 lines 27-31, as follows:

For quantification of mouse GAPDH mRNA the following primers and probes were designed: Sense primer 5'aaggctgtgggcaaggtcatc 3' (SEQ ID NO: 733)(0.3 μ M final concentration),
antisense primer 5' gtcagatccacgacggacacatt (SEQ ID NO: 734) (0.6 μ M final concentration),
TaqMan probe 5' FAM-gaagctcactggcatggcatggccttccgtgttc-TAMRA 3' (SEQ ID NO: 735) (0.2 μ M final concentration).

Please amend the paragraph on page 65 lines 7-13, as follows:

Northern blot analysis was carried out by procedures well known in the art essentially as described in Current Protocols in Molecular Biology, John Wiley & Sons.
The hybridisation probe was obtained by PCR-amplification of a 373 bp fragment from 1 μ l cDNA obtained by reverse transcription PCR. The reaction was carried out using primers 5' agcacaagccattctaagtcattg 3' (SEQ ID NO: 736) (forward) and 5' tccatcatcttacgccagacttc 3' (SEQ ID NO: 737) (reverse) at 0,5 μ M final concentration each, 200 nM each dNTP, 1,5 mM $MgCl_2$ and Platinum Taq DNA polymerase (Invitrogen cat. no. 10966-018).

Please amend the paragraph on page 66 lines 6-11, as follows:

Equality of RNA sample loading was assessed by stripping the blot in 0,5% SDS in H_2O at 85°C and reprobing with a labelled GAPDH (glyceraldehyde-3-phosphate dehydrogenase)

probe obtained essentially as described above using the primers 5' aacggatttggtcgatt 3' (**SEQ ID NO: 739**) (forward) and 5' taagcagttggtggtgca 3' (**SEQ ID NO: 740**) (reverse). See figure 2 and 3. Intensity was monitored with phosphorimager Biorad, FX-scanner (see below). The tested oligomeric compounds are presented in Example 10.

Please amend Table 1 starting on page 67, as follows:

Table 1 Oligomeric compounds of the invention

Oligomeric compounds were evaluated for their potential to knockdown Survivin mRNA in 15PC3 cells. The data are presented as percentage downregulation relative to mock transfected cells. Transcript steady state was monitored by Real-time PCR and normalised to the GAPDH transcript steady state. Note that all LNA C are 5'-Methyl-Cytosine.

Target site	SEQ ID NO.	Oligomeric compound Sequence 5'-3'	Seq ID+ Design	Specific design of Oligomeric compound Capital letters β -D-oxy-LNA S= phosphorthioate O=O-P(O) ₂ -O- Small letters DNA sugar	% Inhibition at 25 nM	% Inhibition at 5 nM oligo
172	2	GCAGTGGATGAAGCCA	2A 147	G _s C _s A _s G _s t _s g _s g _s a _s t _s g _s a _s a _s G _s C _s C _s A	85	44
			2B 148	G _s C _s A _s G _s t _s g _s g _s a _s t _s g _s a _s a _s G _s C _s C _s a	91	
			2C 149	G ₀ C ₀ A ₀ G ₀ t _s g _s g _s a _s t _s g _s a _s a _s G ₀ C ₀ C ₀ A		
			2D 150	g _s c _s a _s g _s t _s g _s g _s a _s t _s g _s a _s a _s g _s c _s c _s a		
198	3	GCCAAGTCTGGCTCGT	3A 151	G _s C _s C _s A _s a _s g _s t _s c _s t _s g _s g _s c _s T _s C _s G _s T	49	
			3B 152	G _s C _s C _s A _s a _s g _s t _s c _s t _s g _s g _s c _s T _s C _s G _s t		
			3C 153	G ₀ C ₀ C ₀ A ₀ a _s g _s t _s c _s t _s g _s g _s c _s T ₀ C ₀ G ₀ T		
			3D 154	g _s c _s c _s a _s a _s g _s t _s c _s t _s g _s g _s c _s t _s c _s g _s t		
206	4	AACACTGGGCCAAGTC	4A 155	A _s A _s C _s A _s c _s t _s g _s g _s g _s c _s c _s a _s A _s G _s T _s C	74	

			<u>4B</u> <u>156</u>	$A_s A_s C_s A_s C_s t_s g_s g_s g_s c_s c_s a_s A_s G_s T_s c$	91	
			<u>4C</u> <u>157</u>	$A_O A_O C_O A_O c_s t_s g_s g_s g_s c_s c_s a_s A_O G_O$ $T_O C$		
			<u>4D</u> <u>158</u>	$a_s a_s c_s a_s c_s t_s g_s g_s g_s c_s c_s a_s a_s g_s t_s c$		
214	5	GCAGAAGAAACACTGG	<u>5A</u> <u>159</u>	$G_s C_s A_s G_s a_s a_s g_s a_s a_s c_s a_s C_s T_s G_s G$	67	
			<u>5B</u> <u>160</u>	$G_s C_s A_s G_s a_s a_s g_s a_s a_s c_s a_s C_s T_s G_s g$		
			<u>5C</u> <u>161</u>	$G_O C_O A_O G_O a_s a_s g_s a_s a_s c_s a_s C_O T_O$ $G_O G$		
			<u>5D</u> <u>162</u>	$g_s c_s a_s g_s a_s a_s g_s a_s a_s c_s a_s c_s t_s g_s g$		
216	6	AAGCAGAAGAAACACT	<u>6A</u> <u>163</u>	$A_s A_s G_s C_s a_s g_s a_s a_s g_s a_s a_s C_s A_s C_s T$	88	63
			<u>6B</u> <u>164</u>	$A_s A_s G_s C_s a_s g_s a_s a_s g_s a_s a_s C_s A_s C_s t$	79	
			<u>6C</u> <u>165</u>	$A_O A_O G_O C_O a_s g_s a_s a_s g_s a_s a_s C_O A_O$ $C_O T$		
			<u>6D</u> <u>166</u>	$a_s a_s g_s c_s a_s g_s a_s a_s g_s a_s a_s c_s a_s c_s t_s$		
238	7	CTCCCAGCCTTCCAGC	<u>7A</u> <u>167</u>	$C_s T_s C_s C_s c_s a_s g_s c_s c_s t_s c_s C_s A_s G_s C$	26	
			<u>7B</u> <u>168</u>	$C_s T_s C_s C_s c_s a_s g_s c_s c_s t_s c_s C_s A_s G_s c$		
			<u>7C</u> <u>169</u>	$C_O T_O C_O C_O c_s a_s g_s c_s c_s t_s c_s C_O A_O G$ O_C		
			<u>7D</u> <u>170</u>	$c_s t_s c_s c_s c_s a_s g_s c_s c_s t_s c_s c_s a_s g_s c$		
403	8	TTCTTTCTTCTTATTG	<u>8A</u> <u>171</u>	$T_s T_s C_s T_s t_s c_s t_s t_s c_s t_s t_s A_s T_s T_s G$	62	
			<u>8B</u> <u>172</u>	$T_s T_s C_s T_s t_s c_s t_s t_s c_s t_s t_s A_s T_s T_s g$		
			<u>8C</u> <u>173</u>	$T_O T_O C_O T_O t_s c_s t_s t_s c_s t_s t_s A_O T_O T_O$ G		
			<u>8D</u> <u>174</u>	$t_s t_s c_s t_s t_s c_s t_s t_s c_s t_s t_s a_s t_s g$		
491	9	TGGGACCAGGCAGCTC	<u>9A</u>	$T_s G_s G_s G_s a_s c_s c_s a_s g_s g_s c_s a_s G_s C_s T_s C$	78	50

			<u>175</u>		
			<u>9B</u> <u>176</u>	$T_s G_s G_s G_s a_s c_s c_s a_s g_s g_s c_s a_s G_s C_s T_s c$	
			<u>9C</u> <u>177</u>	$T_O G_O G_O G_O a_s c_s c_s a_s g_s g_s c_s a_s G_O C_O$ $T_O C$	
			<u>9D</u> <u>178</u>	$t_s t_s c_s t_s t_s t_s c_s t_s t_s c_s t_s a_s t_s t_s g$	
505	10	TGGTGCAGCCACTCTG	<u>10A</u> <u>179</u>	$T_s G_s G_s T_s g_s c_s a_s g_s c_s c_s a_s c_s T_s C_s T_s G$	56
			<u>10B</u> <u>180</u>	$T_s G_s G_s T_s g_s c_s a_s g_s c_s c_s a_s c_s T_s C_s T_s g$	
			<u>10C</u> <u>181</u>	$T_O G_O G_O T_O g_s c_s a_s g_s c_s c_s a_s c_s T_O C_O$ $T_O G$	
			<u>10D</u> <u>182</u>	$t_s g_s g_s t_s g_s c_s a_s g_s c_s c_s a_s c_s t_s c_s t_s g$	
521	11	GAATAAACCTGGAAG	<u>11A</u> <u>183</u>	$G_s A_s A_s T_s a_s a_s a_s c_s c_s t_s g_s G_s A_s A_s G$	58
			<u>11B</u> <u>184</u>	$G_s A_s A_s T_s a_s a_s a_s c_s c_s t_s g_s G_s A_s A_s g$	
			<u>11C</u> <u>185</u>	$G_O A_O A_O T_O a_s a_s a_s c_s c_s t_s g_s G_O A_O$ $A_O G$	
			<u>11D</u> <u>186</u>	$g_s a_s a_s t_s a_s a_s a_s c_s c_s t_s g_s g_s a_s a_s g$	
531	12	TGGCACCAGGGAATAA	<u>12A</u> <u>187</u>	$T_s G_s G_s C_s a_s c_s c_s a_s g_s g_s g_s a_s A_s T_s A_s A$	44
			<u>12B</u> <u>188</u>	$T_s G_s G_s C_s a_s c_s c_s a_s g_s g_s g_s a_s A_s T_s A_s a$	
			<u>12C</u> <u>189</u>	$T_O G_O G_O C_O a_s c_s c_s a_s g_s g_s g_s a_s A_O T_O$ $A_O A$	
			<u>12D</u> <u>190</u>	$t_s g_s g_s c_s a_s c_s c_s a_s g_s g_s g_s a_s t_s a_s a$	
566	13	CTAAGACATTGCTAAG	<u>13A</u> <u>191</u>	$C_s T_s A_s A_s g_s a_s c_s a_s t_s t_s g_s c_s T_s A_s A_s G$	78
			<u>13B</u> <u>192</u>	$C_s T_s A_s A_s g_s a_s c_s a_s t_s t_s g_s c_s T_s A_s A_s g$	
			<u>13C</u> <u>193</u>	$C_O T_O A_O A_O g_s a_s c_s a_s t_s t_s g_s c_s T_O A_O A$ O_G	
			<u>13D</u> <u>194</u>	$c_s t_s a_s a_s g_s a_s c_s a_s t_s t_s g_s c_s t_s a_s a_s g$	

579	14	TTGATCTCCTTTCCTA	14A <u>195</u>	T _s T _s G _s A _s t _s c _s t _s c _s c _s t _s t _s C _s C _s T _s A	73	
			14B <u>196</u>	T _s T _s G _s A _s t _s c _s t _s c _s c _s t _s t _s C _s C _s T _s a		
			14C <u>197</u>	T _O T _O G _O A _O t _s c _s t _s c _s c _s t _s t _s C _O C _O T _O A		
			14D <u>198</u>	t _s t _s g _s a _s t _s c _s t _s c _s c _s t _s t _s c _s c _s t _s a _s		
608	15	GCACAGTTGAAACATC	15A <u>199</u>	G _s C _s A _s C _s a _s g _s t _s g _s a _s a _s C _s A _s T _s C	96	93
			15B <u>200</u>	G _s C _s A _s C _s a _s g _s t _s g _s a _s a _s C _s A _s T _s c	89	79
			15C <u>201</u>	G _O C _O A _O C _O a _s g _s t _s g _s a _s a _s C _O A _O T _O C		
			15D <u>202</u>	g _s c _s a _s c _s a _s g _s t _s g _s a _s a _s c _s a _s t _s c		
			15E <u>203</u>	G _s C _s A _s C _s a _s g _s t _s g _s a _s a _s C _s A _s T _s c	83	78
1	16	GATTCAAATCTGGCGG	16A <u>204</u>	G _s A _s T _s T _s c _s a _s a _s a _s t _s c _s t _s g _s G _s C _s G _s G		
			16B <u>205</u>	G _s A _s T _s T _s c _s a _s a _s a _s t _s c _s t _s g _s G _s C _s G _s g		
			16C <u>206</u>	G _O A _O T _O T _O c _s a _s a _s a _s t _s c _s t _s g _s G _O C _O G _O G		
			16D <u>207</u>	g _s a _s t _s t _s c _s a _s a _s a _s t _s c _s t _s g _s g _s c _s g _s g		
17	17	TGCCAACGGGTCCCGC	17A <u>208</u>	T _s G _s C _s C _s a _s a _s c _s g _s g _s g _s t _s c _s C _s C _s G _s C		
			17B <u>209</u>	T _s G _s C _s C _s a _s a _s c _s g _s g _s g _s t _s c _s C _s C _s G _s c		
			17C <u>210</u>	T _O G _O C _O C _O a _s a _s c _s g _s g _s g _s t _s c _s C _O C _O G _O C		
			17D <u>211</u>	t _s g _s c _s c _s a _s a _s c _s g _s g _s g _s t _s c _s c _s c _s g _s c		
33	18	CCGCCGCCGCCACCTC	18A <u>212</u>	C _s C _s G _s C _s c _s g _s c _s c _s g _s c _s a _s C _s C _s T _s C		
			18B <u>213</u>	C _s C _s G _s C _s c _s g _s c _s c _s g _s c _s a _s C _s C _s T _s c		

			18C <u>214</u>	C ₀ C ₀ G ₀ C ₀ C ₀ g ₀ c ₀ s ₀ c ₀ g ₀ s ₀ c ₀ s ₀ a ₀ C ₀ C ₀ T ₀ C
			18D <u>215</u>	c ₀ s ₀ c ₀ g ₀ s ₀ c ₀ s ₀ g ₀ s ₀ c ₀ s ₀ a ₀ c ₀ s ₀ t ₀ c
49	19	CGTCGGGGCACCCATG	19A <u>216</u>	C _s G _s T _s C _s g _s g _s g _s g _s c _s a _s c _s c _s C _s A _s T _s G
			19B <u>217</u>	C _s G _s T _s C _s g _s g _s g _s g _s c _s a _s c _s c _s C _s A _s T _s g
			19C <u>218</u>	C ₀ G ₀ T ₀ C ₀ g ₀ g ₀ g ₀ g ₀ c ₀ a ₀ c ₀ s ₀ C ₀ A ₀ T ₀ G
			19D <u>219</u>	c _s g _s t _s c _s g _s g _s g _s g _s c _s a _s c _s c _s a _s t _s g
65	20	GCCAGGCAGGGGGCAA	20A <u>220</u>	G _s C _s C _s A _s g _s g _s c _s a _s g _s g _s g _s g _s G _s C _s A _s A
			20B <u>221</u>	G _s C _s C _s A _s g _s g _s c _s a _s g _s g _s g _s g _s G _s C _s A _s a
			20C <u>222</u>	G ₀ C ₀ C ₀ A ₀ g ₀ g ₀ c ₀ a ₀ g ₀ g ₀ g ₀ g ₀ G ₀ C ₀ A ₀ A
			20D <u>223</u>	g _s c _s c _s a _s g _s g _s c _s a _s g _s g _s g _s g _s c _s a _s a
81	21	TCCTTGAGAAAGGGCT	21A <u>224</u>	T _s C _s C _s T _s t _s g _s a _s g _s a _s a _s a _s g _s G _s G _s C _s T
			21B <u>225</u>	T _s C _s C _s T _s t _s g _s a _s g _s a _s a _s a _s g _s G _s G _s C _s t
			21C <u>226</u>	T ₀ C ₀ C ₀ T ₀ t ₀ g ₀ a ₀ g ₀ a ₀ a ₀ a ₀ g ₀ G ₀ G ₀ C ₀ T
			21D <u>227</u>	t _s c _s c _s t _s t _s g _s a _s g _s a _s a _s a _s g _s g _s c _s t
97	22	TGTAGAGATGCGGTGG	22A <u>228</u>	T _s G _s T _s A _s g _s a _s g _s a _s t _s g _s c _s g _s G _s T _s G _s G
			22B <u>229</u>	T _s G _s T _s A _s g _s a _s g _s a _s t _s g _s c _s g _s G _s T _s G _s g
			22C <u>230</u>	T ₀ G ₀ T ₀ A ₀ g ₀ a ₀ g ₀ a ₀ t ₀ g ₀ c ₀ g ₀ G ₀ T ₀ G ₀ G
			22D <u>231</u>	t _s g _s t _s a _s g _s a _s g _s a _s t _s g _s c _s g _s g _s t _s g _s g
113	23	AGGGCCAGTTCTTGAA	23A <u>232</u>	A _s G _s G _s G _s c _s c _s a _s g _s t _s c _s t _s T _s G _s A _s A

			<u>23B</u> <u>233</u>	A _s G _s G _s G _s c _s c _s a _s g _s t _s t _s c _s t _s T _s G _s A _s a
			<u>23C</u> <u>234</u>	A _O G _O G _O G _O c _s c _s a _s g _s t _s t _s c _s t _s T _O G _O A _O A
			<u>23D</u> <u>235</u>	a _s g _s g _s g _s c _s c _s a _s g _s t _s t _s c _s t _s t _s g _s a _s a
129	24	GCGCAGCCCTCCAAGA	<u>24A</u> <u>236</u>	G _s C _s G _s C _s a _s g _s c _s c _s c _s t _s c _s c _s A _s A _s G _s A
			<u>24B</u> <u>237</u>	G _s C _s G _s C _s a _s g _s c _s c _s c _s t _s c _s c _s A _s A _s G _s a
			<u>24C</u> <u>238</u>	G _O C _O G _O C _O a _s g _s c _s c _s c _s t _s c _s c _s A _O A _O G _O A
			<u>24D</u> <u>239</u>	g _s c _s g _s c _s a _s g _s c _s c _s c _s t _s c _s c _s a _s a _s g _s a
145	25	CCGCTCCGGGGTGCAG	<u>25A</u> <u>240</u>	C _s C _s G _s C _s t _s c _s c _s g _s g _s g _s g _s t _s G _s C _s A _s G
			<u>25B</u> <u>241</u>	C _s C _s G _s C _s t _s c _s c _s g _s g _s g _s g _s t _s G _s C _s A _s g
			<u>25C</u> <u>242</u>	C _O C _O G _O C _O t _s c _s c _s g _s g _s g _s g _s t _s G _O C _O A _O G
			<u>25D</u> <u>243</u>	c _s c _s g _s c _s t _s c _s c _s g _s g _s g _s g _s t _s g _s c _s a _s g
161	26	AGCCAGCCTCGGCCAT	<u>26A</u> <u>244</u>	A _s G _s C _s C _s a _s g _s c _s c _s t _s c _s g _s g _s C _s C _s A _s T
			<u>26B</u> <u>245</u>	A _s G _s C _s C _s a _s g _s c _s c _s t _s c _s g _s g _s C _s C _s A _s t
			<u>26C</u> <u>246</u>	A _O G _O C _O C _O a _s g _s c _s c _s t _s c _s g _s g _s C _O C _O A _O T
			<u>26D</u> <u>247</u>	a _s g _s c _s c _s a _s g _s c _s c _s t _s c _s g _s g _s c _s c _s a _s t
177	27	GTGGGGCAGTGGATGA	<u>27A</u> <u>248</u>	G _s T _s G _s G _s g _s g _s c _s a _s g _s t _s g _s g _s A _s T _s G _s A
			<u>27B</u> <u>249</u>	G _s T _s G _s G _s g _s g _s c _s a _s g _s t _s g _s g _s A _s T _s G _s a
			<u>27C</u> <u>250</u>	G _O T _O G _O G _O g _s g _s c _s a _s g _s t _s g _s g _s A _O T _O G _O A
			<u>27D</u> <u>251</u>	g _s t _s g _s g _s g _s c _s a _s g _s t _s g _s g _s a _s t _s g _s a

193	28	GTCTGGCTCGTTCTCA	<u>28A</u> <u>252</u>	$G_s T_s C_s T_s g_s g_s c_s t_s c_s g_s t_s t_s C_s T_s C_s A$
			<u>28B</u> <u>253</u>	$G_s T_s C_s T_s g_s g_s c_s t_s c_s g_s t_s t_s C_s T_s C_s a$
			<u>28C</u> <u>254</u>	$G_O T_O C_O T_O g_s g_s c_s t_s c_s g_s t_s t_s C_O T_O C_O A$
			<u>28D</u> <u>255</u>	$g_s t_s c_s t_s g_s g_s c_s t_s c_s g_s t_s t_s c_s t_s c_s a$
209	29	AGAAACACTGGGCCAA	<u>29A</u> <u>256</u>	$A_s G_s A_s A_s a_s c_s a_s c_s t_s g_s g_s g_s C_s C_s A_s A$
			<u>29B</u> <u>257</u>	$A_s G_s A_s A_s a_s c_s a_s c_s t_s g_s g_s g_s C_s C_s A_s a$
			<u>29C</u> <u>258</u>	$A_O G_O A_O A_O a_s c_s a_s c_s t_s g_s g_s g_s C_O C_O A_O A$
			<u>29D</u> <u>259</u>	$a_s g_s a_s a_s a_s c_s a_s c_s t_s g_s g_s g_s c_s c_s a_s a$
225	30	AGCTCCTTGAAGCAGA	<u>30A</u> <u>260</u>	$A_s G_s C_s T_s c_s c_s t_s t_s g_s a_s a_s g_s C_s A_s G_s A$
			<u>30B</u> <u>261</u>	$A_s G_s C_s T_s c_s c_s t_s t_s g_s a_s a_s g_s C_s A_s G_s a$
			<u>30C</u> <u>262</u>	$A_O G_O C_O T_O c_s c_s t_s t_s g_s a_s a_s g_s C_O A_O G_O A$
			<u>30D</u> <u>263</u>	$a_s g_s c_s t_s c_s c_s t_s t_s g_s a_s a_s g_s c_s a_s g_s a$
241	31	TGGCTCCCAGCCTTCC	<u>31A</u> <u>264</u>	$T_s G_s G_s C_s t_s c_s c_s c_s a_s g_s c_s c_s T_s T_s C_s C$
			<u>31B</u> <u>265</u>	$T_s G_s G_s C_s t_s c_s c_s c_s a_s g_s c_s c_s T_s T_s C_s c$
			<u>31C</u> <u>266</u>	$T_O G_O G_O C_O t_s c_s c_s c_s a_s g_s c_s c_s T_O T_O C_O C$
			<u>31D</u> <u>267</u>	$t_s g_s g_s c_s t_s c_s c_s c_s a_s g_s c_s c_s t_s t_s c_s c$
257	32	CTATGGGGTCGTCATC	<u>32A</u> <u>268</u>	$C_s T_s A_s T_s g_s g_s g_s g_s t_s c_s g_s t_s C_s A_s T_s C$
			<u>32B</u> <u>269</u>	$C_s T_s A_s T_s g_s g_s g_s g_s t_s c_s g_s t_s C_s A_s T_s c$
			<u>32C</u> <u>270</u>	$C_O T_O A_O T_O g_s g_s g_s g_s t_s c_s g_s t_s C_O A_O T_O C$

			<u>32D</u> <u>271</u>	c _s t _s a _s t _s g _s g _s g _s g _s t _s c _s g _s t _s c _s a _s t _s c
273	33	TGCTTTTTATGTTCT	<u>33A</u> <u>272</u>	T _s G _s C _s T _s t _s t _s t _s a _s t _s g _s t _s T _s C _s C _s T
			<u>33B</u> <u>273</u>	T _s G _s C _s T _s t _s t _s t _s a _s t _s g _s t _s T _s C _s C _s t
			<u>33C</u> <u>274</u>	T ₀ G ₀ C ₀ T ₀ t _s t _s t _s a _s t _s g _s t _s T ₀ C ₀ C ₀ O ₀ T
			<u>33D</u> <u>275</u>	t _s g _s c _s t _s t _s t _s t _s a _s t _s g _s t _s c _s c _s t
289	34	AGCGCAACCGGACGAA	<u>34A</u> <u>276</u>	A _s G _s C _s G _s c _s a _s a _s c _s c _s g _s g _s a _s C _s G _s A _s A
			<u>34B</u> <u>277</u>	A _s G _s C _s G _s c _s a _s a _s c _s c _s g _s g _s a _s C _s G _s A _s a
			<u>34C</u> <u>278</u>	A ₀ G ₀ C ₀ G ₀ c _s a _s a _s c _s c _s g _s g _s a _s C ₀ G ₀ A ₀ A
			<u>34D</u> <u>279</u>	a _s g _s c _s g _s c _s a _s a _s c _s c _s g _s g _s a _s c _s g _s a _s a
305	35	TCTTGACAGAAAGGAA	<u>35A</u> <u>280</u>	T _s C _s T _s T _s g _s a _s c _s a _s g _s a _s a _s G _s G _s A _s A
			<u>35B</u> <u>281</u>	T _s C _s T _s T _s g _s a _s c _s a _s g _s a _s a _s G _s G _s A _s a
			<u>35C</u> <u>282</u>	T ₀ C ₀ T ₀ T ₀ g _s a _s c _s a _s g _s a _s a _s G ₀ G ₀ A ₀ A
			<u>35D</u> <u>283</u>	t _s c _s t _s t _s g _s a _s c _s a _s g _s a _s a _s g _s g _s a _s a
321	36	AATTCTTCAAAGGCT	<u>36A</u> <u>284</u>	A _s A _s T _s T _s c _s t _s t _s c _s a _s a _s c _s T _s G _s C _s T
			<u>36B</u> <u>285</u>	A _s A _s T _s T _s c _s t _s t _s c _s a _s a _s c _s T _s G _s C _s t
			<u>36C</u> <u>286</u>	A ₀ A ₀ T ₀ T ₀ c _s t _s t _s c _s a _s a _s c _s T ₀ G ₀ C ₀ O ₀ T
			<u>36D</u> <u>287</u>	a _s a _s t _s t _s c _s t _s t _s c _s a _s a _s c _s t _s g _s c _s t
337	37	AAATTCACCAAGGGTT	<u>37A</u> <u>288</u>	A _s A _s A _s T _s t _s c _s a _s c _s c _s a _s a _s G _s G _s G _s T _s T
			<u>37B</u> <u>289</u>	A _s A _s A _s T _s t _s c _s a _s c _s c _s a _s a _s G _s G _s G _s T _s t

			<u>37C</u> <u>290</u>	A _O A _O A _O T _O t _s c _s a _s c _s c _s a _s a _s g _s G _O G _O T _O T
			<u>37D</u> <u>291</u>	a _s a _s a _s t _s t _s c _s a _s c _s c _s a _s a _s g _s g _s g _s t _s t
353	38	CTCTGTCCAGTTTCAA	<u>38A</u> <u>292</u>	C _s T _s C _s T _s g _s t _s c _s c _s a _s g _s t _s t _s T _s C _s A _s A
			<u>38B</u> <u>293</u>	C _s T _s C _s T _s g _s t _s c _s c _s a _s g _s t _s t _s T _s C _s A _s a
			<u>38C</u> <u>294</u>	C _O T _O C _O T _O g _s t _s c _s c _s a _s g _s t _s t _s T _O C _O A O _A
			<u>38D</u> <u>295</u>	c _s t _s c _s t _s g _s t _s c _s c _s a _s g _s t _s t _s c _s a _s a
369	39	TTGTTCTTGGCTCTTT	<u>39A</u> <u>296</u>	T _s T _s G _s T _s t _s c _s t _s t _s g _s g _s c _s t _s C _s T _s T _s T
			<u>39B</u> <u>297</u>	T _s T _s G _s T _s t _s c _s t _s t _s g _s g _s c _s t _s C _s T _s T _s t
			<u>39C</u> <u>298</u>	T _O T _O G _O T _O t _s c _s t _s t _s g _s g _s c _s t _s C _O T _O T O _T
			<u>39D</u> <u>299</u>	t _s t _s g _s t _s t _s c _s t _s t _s g _s g _s c _s t _s c _s t _s t
385	40	GGTTTCCTTTGCAATT	<u>40A</u> <u>300</u>	G _s G _s T _s T _s t _s c _s c _s t _s t _s g _s c _s A _s A _s T _s T
			<u>40B</u> <u>301</u>	G _s G _s T _s T _s t _s c _s c _s t _s t _s g _s c _s A _s A _s T _s t
			<u>40C</u> <u>302</u>	G _O G _O T _O T _O t _s c _s c _s t _s t _s g _s c _s A _O A _O T O _T
			<u>40D</u> <u>303</u>	g _s g _s t _s t _s c _s c _s t _s t _s g _s c _s a _s a _s t _s t
401	41	CTTTCTTCTTATTGTT	<u>41A</u> <u>304</u>	C _s T _s T _s T _s c _s t _s t _s c _s t _s a _s t _s T _s G _s T _s T
			<u>41B</u> <u>305</u>	C _s T _s T _s T _s c _s t _s t _s c _s t _s a _s t _s T _s G _s T _s t
			<u>41C</u> <u>306</u>	C _O T _O T _O T _O c _s t _s c _s t _s a _s t _s T _O G _O T O _T
			<u>41D</u> <u>307</u>	c _s t _s t _s c _s t _s c _s t _s a _s t _s g _s t _s t
417	42	GCAGTTTCCTCAAATT	<u>42A</u> <u>308</u>	G _s C _s A _s G _s t _s t _s c _s c _s t _s c _s a _s A _s A _s T _s T

			<u>42B</u> <u>309</u>	$G_s C_s A_s G_s t_s t_s c_s c_s t_s c_s a_s A_s A_s T_s t$
			<u>42C</u> <u>310</u>	$G_O C_O A_O G_O t_s t_s c_s c_s t_s c_s a_s A_O A_O T_O T$
			<u>42D</u> <u>311</u>	$g_s c_s a_s g_s t_s t_s c_s c_s t_s c_s a_s a_s t_s t$
433	43	ACGGCGCACTTTCTTC	<u>43A</u> <u>312</u>	$A_s C_s G_s G_s c_s g_s c_s a_s c_s t_s t_s C_s T_s T_s C$
			<u>43B</u> <u>313</u>	$A_s C_s G_s G_s c_s g_s c_s a_s c_s t_s t_s C_s T_s T_s t$
			<u>43C</u> <u>314</u>	$A_O C_O G_O G_O c_s g_s c_s a_s c_s t_s t_s C_O T_O T_O C$
			<u>43D</u> <u>315</u>	$a_s c_s g_s g_s c_s g_s c_s a_s c_s t_s t_s c_s t_s c$
449	44	CCAGCTGCTCGATGGC	<u>44A</u> <u>316</u>	$C_s C_s A_s G_s c_s t_s g_s c_s t_s c_s g_s a_s T_s G_s G_s C$
			<u>44B</u> <u>317</u>	$C_s C_s A_s G_s c_s t_s g_s c_s t_s c_s g_s a_s T_s G_s G_s t$
			<u>44C</u> <u>318</u>	$C_O C_O A_O G_O c_s t_s g_s c_s t_s c_s g_s a_s T_O G_O G_O C$
			<u>44D</u> <u>319</u>	$c_s c_s a_s g_s c_s t_s g_s c_s t_s c_s g_s a_s t_s g_s g_s c$
465	45	CCTCAATCCATGGCAG	<u>45A</u> <u>320</u>	$C_s C_s T_s C_s a_s a_s t_s c_s c_s a_s t_s g_s G_s C_s A_s G$
			<u>45B</u> <u>321</u>	$C_s C_s T_s C_s a_s a_s t_s c_s c_s a_s t_s g_s G_s C_s A_s g$
			<u>45C</u> <u>322</u>	$C_O C_O T_O C_O a_s a_s t_s c_s c_s a_s t_s g_s G_O C_O A_O G$
			<u>45D</u> <u>323</u>	$c_s c_s t_s c_s a_s a_s t_s c_s c_s a_s t_s g_s g_s c_s a_s g$
481	46	CAGCTCCGCCAGAGG	<u>46A</u> <u>324</u>	$C_s A_s G_s C_s t_s c_s c_s g_s g_s c_s c_s a_s G_s A_s G_s G$
			<u>46B</u> <u>325</u>	$C_s A_s G_s C_s t_s c_s c_s g_s g_s c_s c_s a_s G_s A_s G_s g$
			<u>46C</u> <u>326</u>	$C_O A_O G_O C_O t_s c_s c_s g_s g_s c_s c_s a_s G_O A_O G_O G$
			<u>46D</u> <u>327</u>	$c_s a_s g_s c_s t_s c_s c_s g_s g_s c_s c_s a_s g_s a_s g_s g$

497	47	CCACTCTGGGACCAGG	47A 328	C _s C _s A _s C _s t _s c _s t _s g _s g _s g _s a _s c _s C _s A _s G _s G
			47B 329	C _s C _s A _s C _s t _s c _s t _s g _s g _s g _s a _s c _s C _s A _s G _s g
			47C 330	C ₀ C ₀ A ₀ C ₀ t _s c _s t _s g _s g _s g _s a _s c _s C ₀ A ₀ G ₀ G
			47D 331	c _s c _s a _s c _s t _s c _s t _s g _s g _s g _s a _s c _s c _s a _s g _s g
513	48	CCTGGAAGTGGTGCAG	48A 332	C _s C _s T _s G _s g _s a _s a _s g _s t _s g _s g _s t _s G _s C _s A _s G
			48B 333	C _s C _s T _s G _s g _s a _s a _s g _s t _s g _s g _s t _s G _s C _s A _s g
			48C 334	C ₀ C ₀ T ₀ G ₀ g _s a _s a _s g _s t _s g _s g _s t _s G ₀ C ₀ A ₀ G
			48D 335	c _s c _s t _s g _s g _s a _s a _s g _s t _s g _s g _s t _s g _s c _s a _s g
529	49	GCACCAGGGAATAAAC	49A 336	G _s C _s A _s C _s c _s a _s g _s g _s g _s a _s a _s t _s A _s A _s A _s C
			49B 337	G _s C _s A _s C _s c _s a _s g _s g _s g _s a _s a _s t _s A _s A _s A _s c
			49C 338	G ₀ C ₀ A ₀ C ₀ c _s a _s g _s g _s g _s a _s a _s t _s A ₀ A ₀ A ₀ C
			49D 339	g _s c _s a _s c _s c _s a _s g _s g _s g _s a _s a _s t _s a _s a _s a _s c
545	50	CACAGGAAGGCTGGTG	50A 340	C _s A _s C _s A _s g _s g _s a _s a _s g _s c _s t _s G _s G _s T _s G
			50B 341	C _s A _s C _s A _s g _s g _s a _s a _s g _s c _s t _s G _s G _s T _s g
			50C 342	C ₀ A ₀ C ₀ A ₀ g _s g _s a _s a _s g _s c _s t _s G ₀ G ₀ T ₀ G
			50D 343	c _s a _s c _s a _s g _s g _s a _s a _s g _s c _s t _s g _s g _s t _s g
561	51	ACATTGCTAAGGGGCC	51A 344	A _s C _s A _s T _s t _s g _s c _s t _s a _s a _s g _s g _s G _s G _s C _s C
			51B 345	A _s C _s A _s T _s t _s g _s c _s t _s a _s a _s g _s g _s G _s G _s C _s c
			51C 346	A ₀ C ₀ A ₀ T ₀ t _s g _s c _s t _s a _s a _s g _s g _s G ₀ G ₀ C ₀ C

			<u>51D</u> <u>347</u>	$a_s c_s a_s t_s t_s g_s c_s t_s a_s a_s g_s g_s g_s c_s c$
577	52	GATCTCCTTTCCTAAG	<u>52A</u> <u>348</u>	$G_s A_s T_s C_s t_s c_s c_s t_s t_s t_s c_s c_s T_s A_s A_s G$
			<u>52B</u> <u>349</u>	$G_s A_s T_s C_s t_s c_s c_s t_s t_s t_s c_s c_s T_s A_s A_s g$
			<u>52C</u> <u>350</u>	$G_O A_O T_O C_O t_s c_s c_s t_s t_s t_s c_s c_s T_O A_O A_O G$
			<u>52D</u> <u>351</u>	$g_s a_s t_s c_s t_s c_s c_s t_s t_s t_s c_s c_s t_s a_s a_s g$
593	53	CTAATTTGAAAATGTT	<u>53A</u> <u>352</u>	$C_s T_s A_s A_s t_s t_s t_s g_s a_s a_s a_s T_s G_s T_s T$
			<u>53B</u> <u>353</u>	$C_s T_s A_s A_s t_s t_s t_s g_s a_s a_s a_s T_s G_s T_s t$
			<u>53C</u> <u>354</u>	$C_O T_O A_O A_O t_s t_s t_s g_s a_s a_s a_s T_O G_O T_O T$
			<u>53D</u> <u>355</u>	$c_s t_s a_s a_s t_s t_s t_s g_s a_s a_s a_s t_s g_s t_s t$
609	54	AGCACAGTTGAAACAT	<u>54A</u> <u>356</u>	$A_s G_s C_s A_s c_s a_s g_s t_s t_s g_s a_s a_s A_s C_s A_s T$
			<u>54B</u> <u>357</u>	$A_s G_s C_s A_s c_s a_s g_s t_s t_s g_s a_s a_s A_s C_s A_s t$
			<u>54C</u> <u>358</u>	$A_O G_O C_O A_O c_s a_s g_s t_s t_s g_s a_s a_s A_O C_O A_O T$
			<u>54D</u> <u>359</u>	$a_s g_s c_s a_s c_s a_s g_s t_s t_s g_s a_s a_s c_s a_s t$
625	55	TTCAAGACAAAACAGG	<u>55A</u> <u>360</u>	$T_s T_s C_s A_s a_s g_s a_s c_s a_s a_s a_s C_s A_s G_s G$
			<u>55B</u> <u>361</u>	$T_s T_s C_s A_s a_s g_s a_s c_s a_s a_s a_s C_s A_s G_s g$
			<u>55C</u> <u>362</u>	$T_O T_O C_O A_O a_s g_s a_s c_s a_s a_s a_s C_O A_O G_O G$
			<u>55D</u> <u>363</u>	$t_s t_s c_s a_s a_s g_s a_s c_s a_s a_s a_s c_s a_s g_s g$
641	56	CACCTCTGGTGCCACT	<u>56A</u> <u>364</u>	$C_s A_s C_s C_s t_s c_s t_s g_s g_s t_s g_s c_s C_s A_s C_s T$
			<u>56B</u> <u>365</u>	$C_s A_s C_s C_s t_s c_s t_s g_s g_s t_s g_s c_s C_s A_s C_s t$

			<u>56C</u> <u>366</u>	$C_O A_O C_O C_O t_s c_s t_s g_s g_s t_s g_s c_s C_O A_O C_O T$
			<u>56D</u> <u>367</u>	$c_s a_s c_s c_s t_s c_s t_s g_s g_s t_s g_s c_s c_s a_s c_s t$
657	57	GCTGCACAGGCAGAAG	<u>57A</u> <u>368</u>	$G_s C_s T_s G_s c_s a_s c_s a_s g_s g_s c_s a_s G_s A_s A_s G$
			<u>57B</u> <u>369</u>	$G_s C_s T_s G_s c_s a_s c_s a_s g_s g_s c_s a_s G_s A_s A_s g$
			<u>57C</u> <u>370</u>	$G_O C_O T_O G_O c_s a_s c_s a_s g_s g_s c_s a_s G_O A_O A_O G$
			<u>57D</u> <u>371</u>	$g_s c_s t_s g_s c_s a_s c_s a_s g_s g_s c_s a_s g_s a_s a_s g$
673	58	GTTACCAGCAGCACCC	<u>58A</u> <u>372</u>	$G_s T_s T_s A_s c_s c_s a_s g_s c_s a_s g_s c_s A_s C_s C_s C$
			<u>58B</u> <u>373</u>	$G_s T_s T_s A_s c_s c_s a_s g_s c_s a_s g_s c_s A_s C_s C_s c$
			<u>58C</u> <u>374</u>	$G_O T_O T_O A_O c_s c_s a_s g_s c_s a_s g_s c_s A_O C_O C_O C$
			<u>58D</u> <u>375</u>	$g_s t_s t_s a_s c_s c_s a_s g_s c_s a_s g_s c_s a_s c_s c_s c$
689	59	GAGAGAAGCAGCCACT	<u>59A</u> <u>376</u>	$G_s A_s G_s A_s g_s a_s a_s g_s c_s a_s g_s c_s C_s A_s C_s T$
			<u>59B</u> <u>377</u>	$G_s A_s G_s A_s g_s a_s a_s g_s c_s a_s g_s c_s C_s A_s C_s t$
			<u>59C</u> <u>378</u>	$G_O A_O G_O A_O g_s a_s a_s g_s c_s a_s g_s c_s C_O A_O C_O T$
			<u>59D</u> <u>379</u>	$g_s a_s g_s a_s g_s a_s a_s g_s c_s a_s g_s c_s c_s a_s c_s t$
705	60	AAAAAAGAGAGAGAGA	<u>60A</u> <u>380</u>	$A_s A_s A_s A_s a_s a_s g_s a_s g_s a_s g_s a_s G_s A_s G_s A$
			<u>60B</u> <u>381</u>	$A_s A_s A_s A_s a_s a_s g_s a_s g_s a_s g_s a_s G_s A_s G_s a$
			<u>60C</u> <u>382</u>	$A_O A_O A_O A_O a_s a_s g_s a_s g_s a_s g_s a_s G_O A_O G_O A$
			<u>60D</u> <u>383</u>	$a_s a_s a_s a_s a_s g_s a_s g_s a_s g_s a_s g_s a_s g_s a$
721	61	GCAAAAATGAGCCCCC	<u>61A</u> <u>384</u>	$G_s C_s A_s A_s a_s a_s a_s t_s g_s a_s g_s c_s C_s C_s C_s C$

			61B <u>385</u>	G _s C _s A _s A _s a _s a _s a _s t _s g _s a _s g _s c _s C _s C _s C _s c
			61C <u>386</u>	G ₀ C ₀ A ₀ A ₀ a _s a _s a _s t _s g _s a _s g _s c _s C ₀ C ₀ C ₀ C
			61D <u>387</u>	g _s c _s a _s a _s a _s a _s t _s g _s a _s g _s c _s c _s c _s c _s c
737	62	CCCGGGAATCAAAACA	62A <u>388</u>	C _s C _s C _s G _s g _s g _s a _s a _s t _s c _s a _s a _s A _s A _s C _s A
			62B <u>389</u>	C _s C _s C _s G _s g _s g _s a _s a _s t _s c _s a _s a _s A _s A _s C _s a
			62C <u>390</u>	C ₀ C ₀ C ₀ G ₀ g _s g _s a _s a _s t _s c _s a _s a _s A ₀ A ₀ C ₀ A
			62D <u>391</u>	c _s c _s c _s g _s g _s g _s a _s a _s t _s c _s a _s a _s a _s c _s a
753	63	CTTCTCACCTGGTAAG	63A <u>392</u>	C _s T _s T _s C _s t _s c _s a _s c _s c _s t _s g _s g _s T _s A _s A _s G
			63B <u>393</u>	C _s T _s T _s C _s t _s c _s a _s c _s c _s t _s g _s g _s T _s A _s A _s g
			63C <u>394</u>	C ₀ T ₀ T ₀ C ₀ t _s c _s a _s c _s c _s t _s g _s g _s T ₀ A ₀ A ₀ O ₀ G
			63D <u>395</u>	c _s t _s t _s c _s t _s c _s a _s c _s c _s t _s g _s g _s t _s a _s a _s g
769	64	CCTTCTTCCTCCCTCA	64A <u>396</u>	C _s C _s T _s T _s c _s t _s t _s c _s c _s t _s c _s c _s C _s T _s C _s A
			64B <u>397</u>	C _s C _s T _s T _s c _s t _s t _s c _s c _s t _s c _s c _s C _s T _s C _s a
			64C <u>398</u>	C ₀ C ₀ T ₀ T ₀ c _s t _s c _s c _s t _s c _s c _s C ₀ T ₀ C ₀ O ₀ A
			64D <u>399</u>	c _s c _s t _s t _s c _s t _s t _s c _s c _s t _s c _s c _s c _s t _s c _s a
785	65	AGCAAAAGGGACACTG	65A <u>400</u>	A _s G _s C _s A _s a _s a _s a _s g _s g _s g _s a _s c _s A _s C _s T _s G
			65B <u>401</u>	A _s G _s C _s A _s a _s a _s a _s g _s g _s g _s a _s c _s A _s C _s T _s g
			65C <u>402</u>	A ₀ G ₀ C ₀ A ₀ a _s a _s a _s g _s g _s g _s a _s c _s A ₀ C ₀ T ₀ G
			65D <u>403</u>	a _s g _s c _s a _s a _s a _s a _s g _s g _s g _s a _s c _s a _s c _s t _s g

801	66	CAAAGCTGTCAGCTCT	<u>66A</u> <u>404</u>	C _s A _s A _s A _s g _s c _s t _s g _s t _s c _s a _s g _s C _s T _s C _s T
			<u>66B</u> <u>405</u>	C _s A _s A _s A _s g _s c _s t _s g _s t _s c _s a _s g _s C _s T _s C _s t
			<u>66C</u> <u>406</u>	C ₀ A ₀ A ₀ A ₀ g _s c _s t _s g _s t _s c _s a _s g _s C ₀ T ₀ C ₀ T
			<u>66D</u> <u>407</u>	c _s a _s a _s a _s g _s c _s t _s g _s t _s c _s a _s g _s c _s t _s c _s t
817	67	GCTCTGCCCACGCGAA	<u>67A</u> <u>408</u>	G _s C _s T _s C _s t _s g _s c _s c _s c _s a _s c _s g _s C _s G _s A _s A
			<u>67B</u> <u>409</u>	G _s C _s T _s C _s t _s g _s c _s c _s c _s a _s c _s g _s C _s G _s A _s a
			<u>67C</u> <u>410</u>	G ₀ C ₀ T ₀ C ₀ t _s g _s c _s c _s c _s a _s c _s g _s C ₀ G ₀ A ₀ A
			<u>67D</u> <u>411</u>	g _s c _s t _s c _s t _s g _s c _s c _s c _s a _s c _s g _s c _s g _s a _s a
833	68	ACATTCAGGTGGAAG	<u>68A</u> <u>412</u>	A _s C _s A _s T _s t _s c _s a _s c _s t _s g _s t _s g _s G _s A _s A _s G
			<u>68B</u> <u>413</u>	A _s C _s A _s T _s t _s c _s a _s c _s t _s g _s t _s g _s G _s A _s A _s g
			<u>68C</u> <u>414</u>	A ₀ C ₀ A ₀ T ₀ t _s c _s a _s c _s t _s g _s t _s g _s G ₀ A ₀ A O ₀ G
			<u>68D</u> <u>415</u>	a _s c _s a _s t _s c _s a _s c _s t _s g _s t _s g _s a _s a _s g
849	69	AACATGAGGTCCAGAC	<u>69A</u> <u>416</u>	A _s A _s C _s A _s t _s g _s a _s g _s g _s t _s c _s c _s A _s G _s A _s C
			<u>69B</u> <u>417</u>	A _s A _s C _s A _s t _s g _s a _s g _s g _s t _s c _s c _s A _s G _s A _s c
			<u>69C</u> <u>418</u>	A ₀ A ₀ C ₀ A ₀ t _s g _s a _s g _s g _s t _s c _s c _s A ₀ G ₀ A ₀ C
			<u>69D</u> <u>419</u>	a _s a _s c _s a _s t _s g _s a _s g _s g _s t _s c _s c _s a _s g _s a _s c
865	70	CTGTGACAGCCTCAAC	<u>70A</u> <u>420</u>	C _s T _s G _s T _s g _s a _s c _s a _s g _s c _s c _s t _s C _s A _s A _s C
			<u>70B</u> <u>421</u>	C _s T _s G _s T _s g _s a _s c _s a _s g _s c _s c _s t _s C _s A _s A _s c
			<u>70C</u> <u>422</u>	C ₀ T ₀ G ₀ T ₀ g _s a _s c _s a _s g _s c _s c _s t _s C ₀ A ₀ A ₀ C

			<u>70D</u> <u>423</u>	c _s t _s g _s t _s g _s a _s c _s a _s g _s c _s c _s t _s c _s a _s a _s c
881	71	AAGTCCACACTCAGGA	<u>71A</u> <u>424</u>	A _s A _s G _s T _s c _s c _s a _s c _s a _s c _s t _s c _s A _s G _s G _s A
			<u>71B</u> <u>425</u>	A _s A _s G _s T _s c _s c _s a _s c _s a _s c _s t _s c _s A _s G _s G _s a
			<u>71C</u> <u>426</u>	A ₀ A ₀ G ₀ T ₀ c _s c _s a _s c _s a _s c _s t _s c _s A ₀ G ₀ G ₀ A
			<u>71D</u> <u>427</u>	a _s a _s g _s t _s c _s c _s a _s c _s a _s c _s t _s c _s a _s g _s g _s a
897	72	TCAACAGGCACCTGCC	<u>72A</u> <u>428</u>	T _s C _s A _s A _s c _s a _s g _s g _s c _s a _s c _s T _s G _s C _s C
			<u>72B</u> <u>429</u>	T _s C _s A _s A _s c _s a _s g _s g _s c _s a _s c _s T _s G _s C _s c
			<u>72C</u> <u>430</u>	T ₀ C ₀ A ₀ A ₀ c _s a _s g _s g _s c _s a _s c _s T ₀ G ₀ C ₀ C
			<u>72D</u> <u>431</u>	t _s c _s a _s a _s c _s a _s g _s g _s c _s a _s c _s t _s g _s c _s c
913	73	AACCTGCAGCTCAGAT	<u>73A</u> <u>432</u>	A _s A _s C _s C _s t _s g _s c _s a _s g _s c _s t _s c _s A _s G _s A _s T
			<u>73B</u> <u>433</u>	A _s A _s C _s C _s t _s g _s c _s a _s g _s c _s t _s c _s A _s G _s A _s t
			<u>73C</u> <u>434</u>	A ₀ A ₀ C ₀ C ₀ t _s g _s c _s a _s g _s c _s t _s c _s A ₀ G ₀ A ₀ T
			<u>73D</u> <u>435</u>	a _s a _s c _s c _s t _s g _s c _s a _s g _s c _s t _s c _s a _s g _s a _s t
929	74	GGTGTGACAGATAAGG	<u>74A</u> <u>436</u>	G _s G _s T _s G _s t _s g _s a _s c _s a _s g _s a _s t _s A _s A _s G _s G
			<u>74B</u> <u>437</u>	G _s G _s T _s G _s t _s g _s a _s c _s a _s g _s a _s t _s A _s A _s G _s g
			<u>74C</u> <u>438</u>	G ₀ G ₀ T ₀ G ₀ t _s g _s a _s c _s a _s g _s a _s t _s A ₀ A ₀ G ₀ G
			<u>74D</u> <u>439</u>	g _s g _s t _s g _s t _s g _s a _s c _s a _s g _s a _s t _s a _s a _s g _s g
945	75	CCTCTGAGGAGGCACA	<u>75A</u> <u>440</u>	C _s C _s T _s C _s t _s g _s a _s g _s g _s a _s g _s C _s A _s C _s A
			<u>75B</u> <u>441</u>	C _s C _s T _s C _s t _s g _s a _s g _s g _s a _s g _s C _s A _s C _s a

			<u>75C</u> <u>442</u>	C _O C _O T _O C _O t _S g _S a _S g _S g _S a _S g _S g _S C _O A _O C _O A
			<u>75D</u> <u>443</u>	c _S c _S t _S c _S t _S g _S a _S g _S g _S a _S g _S g _S c _S a _S c _S a
961	76	ACAACAAAAAACTGT	<u>76A</u> <u>444</u>	A _S C _S A _S A _S c _S a _S a _S a _S a _S a _S a _S C _S T _S G _S T
			<u>76B</u> <u>445</u>	A _S C _S A _S A _S c _S a _S a _S a _S a _S a _S a _S C _S T _S G _S t
			<u>76C</u> <u>446</u>	A _O C _O A _O A _O c _S a _S a _S a _S a _S a _S a _S C _O T _O G _O T
			<u>76D</u> <u>447</u>	a _S c _S a _S a _S c _S a _S a _S a _S a _S a _S a _S c _S t _S g _S t
977	77	AAAACAAAAAAACACA	<u>77A</u> <u>448</u>	A _S A _S A _S A _S c _S a _S a _S a _S a _S a _S a _S C _S A _S C _S A
			<u>77B</u> <u>449</u>	A _S A _S A _S A _S c _S a _S a _S a _S a _S a _S a _S C _S A _S C _S a
			<u>77C</u> <u>450</u>	A _O A _O A _O A _O c _S a _S a _S a _S a _S a _S a _S C _O A _O C _O A
			<u>77D</u> <u>451</u>	a _S a _S a _S a _S c _S a _S a _S a _S a _S a _S a _S c _S a _S c _S a
993	78	CATCTACCAAAAAAAA	<u>78A</u> <u>452</u>	C _S A _S T _S C _S t _S a _S c _S c _S a _S a _S a _S a _S A _S A _S A _S A
			<u>78B</u> <u>453</u>	C _S A _S T _S C _S t _S a _S c _S c _S a _S a _S a _S a _S A _S A _S A _S a
			<u>78C</u> <u>454</u>	C _O A _O T _O C _O t _S a _S c _S c _S a _S a _S a _S a _S A _O A _O A _O A
			<u>78D</u> <u>455</u>	c _S a _S t _S c _S t _S a _S c _S c _S a _S a _S a _S a _S a _S a
1009	79	TCACACACAAGTCATG	<u>79A</u> <u>456</u>	T _S C _S A _S C _S a _S c _S a _S c _S a _S a _S g _S t _S C _S A _S T _S G
			<u>79B</u> <u>457</u>	T _S C _S A _S C _S a _S c _S a _S c _S a _S a _S g _S t _S C _S A _S T _S g
			<u>79C</u> <u>458</u>	T _O C _O A _O C _O a _S c _S a _S c _S a _S a _S g _S t _S C _O A _O T _O G
			<u>79D</u> <u>459</u>	t _S c _S a _S c _S a _S c _S a _S c _S a _S a _S g _S t _S c _S a _S t _S g
1025	80	TGTCTCCATTCTCTCA	<u>80A</u> <u>460</u>	T _S G _S T _S C _S t _S c _S c _S a _S t _S c _S t _S C _S T _S C _S A

			<u>80B</u> <u>461</u>	T _s G _s T _s C _s t _s c _s c _s a _s t _s c _s t _s C _s T _s C _s a
			<u>80C</u> <u>462</u>	T _O G _O T _O C _O t _s c _s c _s a _s t _s c _s t _s C _O T _O C _O A
			<u>80D</u> <u>463</u>	t _s g _s t _s c _s t _s c _s c _s a _s t _s c _s t _s c _s t _s c _s a
1041	81	GAGGAGCCAGGGACTC	<u>81A</u> <u>464</u>	G _s A _s G _s G _s a _s g _s c _s c _s a _s g _s g _s g _s A _s C _s T _s C
			<u>81B</u> <u>465</u>	G _s A _s G _s G _s a _s g _s c _s c _s a _s g _s g _s g _s A _s C _s T _s c
			<u>81C</u> <u>466</u>	G _O A _O G _O G _O a _s g _s c _s c _s a _s g _s g _s g _s A _O C _O T _O C
			<u>81D</u> <u>467</u>	g _s a _s g _s g _s a _s g _s c _s c _s a _s g _s g _s g _s a _s c _s t _s c
1057	82	ATGTTGTAAACAGTA	<u>82A</u> <u>468</u>	A _s T _s G _s T _s t _s g _s t _s t _s a _s a _s c _s A _s G _s T _s A
			<u>82B</u> <u>469</u>	A _s T _s G _s T _s t _s g _s t _s t _s a _s a _s c _s A _s G _s T _s a
			<u>82C</u> <u>470</u>	A _O T _O G _O T _O t _s g _s t _s t _s a _s a _s c _s A _O G _O T _O A
			<u>82D</u> <u>471</u>	a _s t _s g _s t _s t _s g _s t _s t _s a _s a _s c _s a _s g _s t _s a
1073	83	ACAAAATAAGAAAGCC	<u>83A</u> <u>472</u>	A _s C _s A _s A _s a _s a _s t _s a _s a _s g _s a _s a _s A _s G _s C _s C
			<u>83B</u> <u>473</u>	A _s C _s A _s A _s a _s a _s t _s a _s a _s g _s a _s a _s A _s G _s C _s c
			<u>83C</u> <u>474</u>	A _O C _O A _O A _O a _s a _s t _s a _s a _s g _s a _s a _s A _O G _O C _O C
			<u>83D</u> <u>475</u>	a _s c _s a _s a _s a _s t _s a _s a _s g _s a _s a _s g _s c _s c
1089	84	TGAATTAACAATTCAA	<u>84A</u> <u>476</u>	T _s G _s A _s A _s t _s t _s a _s a _s c _s a _s a _s t _s T _s C _s A _s A
			<u>84B</u> <u>477</u>	T _s G _s A _s A _s t _s t _s a _s a _s c _s a _s a _s t _s T _s C _s A _s a
			<u>84C</u> <u>478</u>	T _O G _O A _O A _O t _s t _s a _s a _s c _s a _s a _s t _s T _O C _O A _O A
			<u>84D</u> <u>479</u>	t _s g _s a _s a _s t _s t _s a _s a _s c _s a _s a _s t _s t _s c _s a _s a

1105	85	AGTTTGTGCTATTCTG	<u>85A</u> <u>480</u>	A _s G _s T _s T _s t _s g _s t _s g _s c _s t _s a _s t _s T _s C _s T _s G
			<u>85B</u> <u>481</u>	A _s G _s T _s T _s t _s g _s t _s g _s c _s t _s a _s t _s T _s C _s T _s g
			<u>85C</u> <u>482</u>	A _O G _O T _O T _O t _s g _s t _s g _s c _s t _s a _s t _s T _O C _O T _O G
			<u>85D</u> <u>483</u>	a _s g _s t _s t _s t _s g _s t _s g _s c _s t _s a _s t _s c _s t _s g
1121	86	GCTTAGTTTTAATTGT	<u>86A</u> <u>484</u>	G _s C _s T _s T _s a _s g _s t _s t _s t _s a _s a _s T _s T _s G _s T
			<u>86B</u> <u>485</u>	G _s C _s T _s T _s a _s g _s t _s t _s t _s a _s a _s T _s T _s G _s t
			<u>86C</u> <u>486</u>	G _O C _O T _O T _O a _s g _s t _s t _s t _s a _s a _s T _O T _O G _O T
			<u>86D</u> <u>487</u>	g _s c _s t _s t _s a _s g _s t _s t _s t _s a _s a _s t _s g _s t
1137	87	CTTAGAATGGCTTTGT	<u>87A</u> <u>488</u>	C _s T _s T _s A _s g _s a _s a _s t _s g _s g _s c _s t _s T _s T _s G _s T
			<u>87B</u> <u>489</u>	C _s T _s T _s A _s g _s a _s a _s t _s g _s g _s c _s t _s T _s T _s G _s t
			<u>87C</u> <u>490</u>	C _O T _O T _O A _O g _s a _s a _s t _s g _s g _s c _s t _s T _O T _O G _O T
			<u>87D</u> <u>491</u>	c _s t _s t _s a _s g _s a _s a _s t _s g _s g _s c _s t _s t _s g _s t
1153	88	CCCGTTTCCCCAATGA	<u>88A</u> <u>492</u>	C _s C _s C _s G _s t _s t _s c _s c _s c _s c _s a _s A _s T _s G _s A
			<u>88B</u> <u>493</u>	C _s C _s C _s G _s t _s t _s c _s c _s c _s c _s a _s A _s T _s G _s a
			<u>88C</u> <u>494</u>	C _O C _O C _O G _O t _s t _s c _s c _s c _s c _s a _s A _O T _O G _O A
			<u>88D</u> <u>495</u>	c _s c _s c _s g _s t _s t _s c _s c _s c _s c _s a _s a _s t _s g _s a
1169	89	TCCACCTGAAGTTCAC	<u>89A</u> <u>496</u>	T _s C _s C _s A _s c _s c _s t _s g _s a _s g _s t _s T _s C _s A _s C
			<u>89B</u> <u>497</u>	T _s C _s C _s A _s c _s c _s t _s g _s a _s g _s t _s T _s C _s A _s c
			<u>89C</u> <u>498</u>	T _O C _O C _O A _O c _s t _s g _s a _s g _s t _s T _O C _O A _O C

			<u>89D</u> <u>499</u>	$t_s c_s c_s a_s c_s c_s t_s g_s a_s a_s g_s t_s c_s a_s c$
1185	90	CTATTCTGTCTCCTCA	<u>90A</u> <u>500</u>	$C_s T_s A_s T_s t_s c_s t_s g_s t_s c_s t_s c_s C_s T_s C_s A$
			<u>90B</u> <u>501</u>	$C_s T_s A_s T_s t_s c_s t_s g_s t_s c_s t_s c_s C_s T_s C_s a$
			<u>90C</u> <u>502</u>	$C_O T_O A_O T_O t_s c_s t_s g_s t_s c_s t_s c_s C_O T_O C_O A$
			<u>90D</u> <u>503</u>	$c_s t_s a_s t_s t_s c_s t_s g_s t_s c_s t_s c_s t_s c_s a$
1201	91	GACGCTTCCTATCACT	<u>91A</u> <u>504</u>	$G_s A_s C_s G_s c_s t_s t_s c_s c_s t_s a_s t_s C_s A_s C_s T$
			<u>91B</u> <u>505</u>	$G_s A_s C_s G_s c_s t_s t_s c_s c_s t_s a_s t_s C_s A_s C_s t$
			<u>91C</u> <u>506</u>	$G_O A_O C_O G_O c_s t_s t_s c_s c_s t_s a_s t_s C_O A_O C_O T$
			<u>91D</u> <u>507</u>	$g_s a_s c_s g_s c_s t_s t_s c_s c_s t_s a_s t_s c_s a_s c_s t$
1217	92	AAAGGAGTATCTGCCA	<u>92A</u> <u>508</u>	$A_s A_s A_s G_s g_s a_s g_s t_s a_s t_s c_s t_s G_s C_s C_s A$
			<u>92B</u> <u>509</u>	$A_s A_s A_s G_s g_s a_s g_s t_s a_s t_s c_s t_s G_s C_s C_s a$
			<u>92C</u> <u>510</u>	$A_O A_O A_O G_O g_s a_s g_s t_s a_s t_s c_s t_s G_O C_O C_O A$
			<u>92D</u> <u>511</u>	$a_s a_s a_s g_s g_s a_s g_s t_s a_s t_s c_s t_s g_s c_s c_s a$
1233	93	TCACACAGCAGTGGCA	<u>93A</u> <u>512</u>	$T_s C_s A_s C_s a_s c_s a_s g_s c_s a_s g_s t_s G_s G_s C_s A$
			<u>93B</u> <u>513</u>	$T_s C_s A_s C_s a_s c_s a_s g_s c_s a_s g_s t_s G_s G_s C_s a$
			<u>93C</u> <u>514</u>	$T_O C_O A_O C_O a_s c_s a_s g_s c_s a_s g_s t_s G_O G_O C_O A$
			<u>93D</u> <u>515</u>	$t_s c_s a_s c_s a_s c_s a_s g_s c_s a_s g_s t_s g_s g_s c_s a$
1249	94	CACTGGGCCTGTCTAA	<u>94A</u> <u>516</u>	$C_s A_s C_s T_s g_s g_s g_s c_s c_s t_s g_s t_s C_s T_s A_s A$
			<u>94B</u> <u>517</u>	$C_s A_s C_s T_s g_s g_s g_s c_s c_s t_s g_s t_s C_s T_s A_s a$

			<u>94C</u> <u>518</u>	$C_O A_O C_O T_O g_S g_S g_S c_S c_S t_S g_S t_S C_O T_O$ $A_O A$
			<u>94D</u> <u>519</u>	$c_S a_S c_S t_S g_S g_S g_S c_S c_S t_S g_S t_S c_S t_S a_S a$
1265	95	CATGTGCCCCGCGGCT	<u>95A</u> <u>520</u>	$C_S A_S T_S G_S t_S g_S c_S c_S c_S c_S g_S c_S G_S G_S C_S T$
			<u>95B</u> <u>521</u>	$C_S A_S T_S G_S t_S g_S c_S c_S c_S c_S g_S c_S G_S G_S C_S t$
			<u>95C</u> <u>522</u>	$C_O A_O T_O G_O t_S g_S c_S c_S c_S c_S g_S c_S G_O G_O$ $C_O T$
			<u>95D</u> <u>523</u>	$c_S a_S t_S g_S t_S g_S c_S c_S c_S c_S g_S c_S g_S g_S c_S t$
1281	96	AGGGAGGAGCGGCCAG	<u>96A</u> <u>524</u>	$A_S G_S G_S G_S a_S g_S g_S a_S g_S c_S g_S g_S C_S C_S A_S G$
			<u>96B</u> <u>525</u>	$A_S G_S G_S G_S a_S g_S g_S a_S g_S c_S g_S g_S C_S C_S A_S g$
			<u>96C</u> <u>526</u>	$A_O G_O G_O G_O a_S g_S g_S a_S g_S c_S g_S g_S C_O C_O$ $A_O G$
			<u>96D</u> <u>527</u>	$a_S g_S g_S g_S a_S g_S g_S a_S g_S c_S g_S g_S c_S c_S a_S g$
1297	97	CCACTGCCTTTTCTG	<u>97A</u> <u>528</u>	$C_S C_S A_S C_S t_S g_S c_S c_S t_S t_S t_S T_S C_S T_S G$
			<u>97B</u> <u>529</u>	$C_S C_S A_S C_S t_S g_S c_S c_S t_S t_S t_S T_S C_S T_S g$
			<u>97C</u> <u>530</u>	$C_O C_O A_O C_O t_S g_S c_S c_S t_S t_S t_S T_O C_O T$ O_G
			<u>97D</u> <u>531</u>	$c_S c_S a_S c_S t_S g_S c_S c_S t_S t_S t_S t_S c_S t_S g$
1313	98	TTAAAAAGGATTTAGG	<u>98A</u> <u>532</u>	$T_S T_S A_S A_S a_S a_S a_S g_S g_S a_S t_S T_S A_S G_S G$
			<u>98B</u> <u>533</u>	$T_S T_S A_S A_S a_S a_S a_S g_S g_S a_S t_S T_S A_S G_S g$
			<u>98C</u> <u>534</u>	$T_O T_O A_O A_O a_S a_S a_S g_S g_S a_S t_S T_O A_O G$ O_G
			<u>98D</u> <u>535</u>	$t_S t_S a_S a_S a_S a_S g_S g_S a_S t_S t_S a_S g_S g$
1329	99	CATCGAGCCAAGTCAT	<u>99A</u> <u>536</u>	$C_S A_S T_S C_S g_S a_S g_S c_S c_S a_S a_S g_S T_S C_S A_S T$

			99B <u>537</u>	C _s A _s T _s C _s g _s a _s g _s c _s c _s a _s a _s g _s T _s C _s A _s t
			99C <u>538</u>	C _O A _O T _O C _O g _s a _s g _s c _s c _s a _s g _s T _O C _O A _O T
			99D <u>539</u>	c _s a _s t _s c _s g _s a _s g _s c _s c _s a _s a _s g _s t _s c _s a _s t
1345	100	AGCCAGTCCCCACAG	100A <u>540</u>	A _s G _s C _s C _s a _s g _s t _s c _s c _s c _s c _s A _s C _s A _s G
			100B <u>541</u>	A _s G _s C _s C _s a _s g _s t _s c _s c _s c _s c _s A _s C _s A _s g
			100C <u>542</u>	A _O G _O C _O C _O a _s g _s t _s c _s c _s c _s c _s A _O C _O A _O G
			100D <u>543</u>	a _s g _s c _s c _s a _s g _s t _s c _s c _s c _s c _s a _s c _s a _s g
1361	101	CGGCCTGCAGCAGCCC	101A <u>544</u>	C _s G _s G _s C _s c _s t _s g _s c _s a _s g _s c _s a _s G _s C _s C _s C
			101B <u>545</u>	C _s G _s G _s C _s c _s t _s g _s c _s a _s g _s c _s a _s G _s C _s C _s c
			101C <u>546</u>	C _O G _O G _O C _O c _s t _s g _s c _s a _s g _s c _s a _s G _O C _O C _O C
			101D <u>547</u>	c _s g _s g _s c _s c _s t _s g _s c _s a _s g _s c _s a _s g _s c _s c
1377	102	TGGGCTGACAGACACA	102A <u>548</u>	T _s G _s G _s G _s c _s t _s g _s a _s c _s a _s g _s a _s C _s A _s C _s A
			102B <u>549</u>	T _s G _s G _s G _s c _s t _s g _s a _s c _s a _s g _s a _s C _s A _s C _s a
			102C <u>550</u>	T _O G _O G _O G _O c _s t _s g _s a _s c _s a _s g _s a _s C _O A _O C _O A
			102D <u>551</u>	t _s g _s g _s g _s c _s t _s g _s a _s c _s a _s g _s a _s c _s a _s c _s a
1393	103	TGACAGATGTGAAGGT	103A <u>552</u>	T _s G _s A _s C _s a _s g _s a _s t _s g _s t _s g _s a _s A _s G _s G _s T

			103B <u>553</u>	T _s G _s A _s C _s a _s g _s a _s t _s g _s t _s g _s a _s A _s G _s G _s t
			103C <u>554</u>	T _O G _O A _O C _O a _s g _s a _s t _s g _s t _s g _s a _s A _O G _O G _O T
			103D <u>555</u>	t _s g _s a _s c _s a _s g _s a _s t _s g _s t _s g _s a _s a _s g _s g _s t
1409	104	CCCCGTGTGGAGAACG	104A <u>556</u>	C _s C _s C _s C _s g _s t _s g _s t _s g _s g _s a _s g _s A _s A _s C _s G
			104B <u>557</u>	C _s C _s C _s C _s g _s t _s g _s t _s g _s g _s a _s g _s A _s A _s C _s g
			1004C <u>558</u>	C _O C _O C _O C _O g _s t _s g _s t _s g _s g _s a _s g _s A _O A _O C _O G
			104D <u>559</u>	c _s c _s c _s c _s g _s t _s g _s t _s g _s g _s a _s g _s a _s a _s c _s g
1425	105	GCGGACTGCGTCTCTC	105A <u>560</u>	G _s C _s G _s G _s a _s c _s t _s g _s c _s g _s t _s c _s T _s C _s T _s C
			105B <u>561</u>	G _s C _s G _s G _s a _s c _s t _s g _s c _s g _s t _s c _s T _s C _s T _s c
			105C <u>562</u>	G _O C _O G _O G _O a _s c _s t _s g _s c _s g _s t _s c _s T _O C _O T _O C
			105D <u>563</u>	g _s c _s g _s g _s a _s c _s t _s g _s c _s g _s t _s c _s t _s c _s t _s c
1441	106	GAAAGCGGGGACCTGG	106A <u>564</u>	G _s A _s A _s A _s g _s c _s g _s g _s g _s g _s a _s c _s C _s T _s G _s G
			106B <u>565</u>	G _s A _s A _s A _s g _s c _s g _s g _s g _s g _s a _s c _s C _s T _s G _s g
			106C <u>566</u>	G _O A _O A _O A _O g _s c _s g _s g _s g _s g _s a _s c _s C _O T _O G _O G
			106D <u>567</u>	g _s a _s a _s a _s g _s c _s g _s g _s g _s g _s a _s c _s c _s t _s g _s g

1457	107	AGCTGCTGCCTCCAAA	107 A <u>568</u>	A _s G _s C _s T _s g _s c _s t _s g _s c _s c _s t _s c _s C _s A _s A _s A
			107B <u>569</u>	A _s G _s C _s T _s g _s c _s t _s g _s c _s c _s t _s c _s C _s A _s A _s a
			107C <u>570</u>	A ₀ G ₀ C ₀ T ₀ g _s c _s t _s g _s c _s c _s t _s c _s C ₀ A ₀ A ₀ A
			107 D <u>571</u>	a _s g _s c _s t _s g _s c _s t _s g _s c _s c _s t _s c _s a _s a _s a
1473	108	ACTTCAGCCCTGCGGG	108 A <u>572</u>	A _s C _s T _s T _s c _s a _s g _s c _s c _s t _s g _s C _s G _s G _s G
			108B <u>573</u>	A _s C _s T _s T _s c _s a _s g _s c _s c _s t _s g _s C _s G _s G _s g
			108C <u>574</u>	A ₀ C ₀ T ₀ T ₀ c _s a _s g _s c _s c _s t _s g _s C ₀ G ₀ G ₀ G
			108 D <u>575</u>	a _s c _s t _s t _s c _s a _s g _s c _s c _s c _s t _s g _s c _s g _s g _s g
1489	109	CATCATCTTACGCCAG	109 A <u>576</u>	C _s A _s T _s C _s a _s t _s c _s t _s a _s c _s g _s C _s C _s A _s G
			109B <u>577</u>	C _s A _s T _s C _s a _s t _s c _s t _s a _s c _s g _s C _s C _s A _s g
			109C <u>578</u>	C ₀ A ₀ T ₀ C ₀ a _s t _s c _s t _s a _s c _s g _s C ₀ C ₀ A O ₀ G
			109 D <u>579</u>	c _s a _s t _s c _s a _s t _s c _s t _s a _s c _s g _s c _s c _s a _s g
1505	110	GAGGGCGAATCAAATC	110 A <u>580</u>	G _s A _s G _s G _s g _s c _s g _s a _s a _s t _s c _s a _s A _s A _s T _s C
			110B <u>581</u>	G _s A _s G _s G _s g _s c _s g _s a _s a _s t _s c _s a _s A _s A _s T _s c
			110C <u>582</u>	G ₀ A ₀ G ₀ G ₀ g _s c _s g _s a _s a _s t _s c _s a _s A ₀ A ₀ T ₀ C
			110 D <u>583</u>	g _s a _s g _s g _s c _s g _s a _s a _s t _s c _s a _s a _s a _s t _s c

1521	111	GCTCTATGACAGGGAG	111 A <u>584</u>	G _s C _s T _s C _s t _s a _s t _s g _s a _s c _s a _s g _s G _s G _s A _s G
			111B <u>585</u>	G _s C _s T _s C _s t _s a _s t _s g _s a _s c _s a _s g _s G _s G _s A _s g
			111C <u>586</u>	G _O C _O T _O C _O t _s a _s t _s g _s a _s c _s a _s g _s G _O G _O A _O G
			111 D <u>587</u>	g _s c _s t _s c _s t _s a _s t _s g _s a _s c _s a _s g _s g _s a _s g
1537	112	AACAATCCACCCTGCA	112 A <u>588</u>	A _s A _s C _s A _s a _s t _s c _s c _s a _s c _s c _s T _s G _s C _s A
			112B <u>589</u>	A _s A _s C _s A _s a _s t _s c _s c _s a _s c _s c _s T _s G _s C _s a
			112C <u>590</u>	A _O A _O C _O A _O a _s t _s c _s c _s a _s c _s c _s T _O G _O C _O A
			112 D <u>591</u>	a _s a _s c _s a _s a _s t _s c _s c _s a _s c _s c _s t _s g _s c _s a
1553	113	TTTCCAGCGAAGCTGT	113 A <u>592</u>	T _s T _s T _s C _s c _s a _s g _s c _s g _s a _s a _s g _s C _s T _s G _s T
			113B <u>593</u>	T _s T _s T _s C _s c _s a _s g _s c _s g _s a _s a _s g _s C _s T _s G _s t
			113C <u>594</u>	T _O T _O T _O C _O c _s a _s g _s c _s g _s a _s a _s g _s C _O T _O G _O T
			113 D <u>595</u>	t _s t _s t _s c _s c _s a _s g _s c _s g _s a _s a _s g _s c _s t _s g _s t
1569	114	AGATGACCTCCAGAGG	114 A <u>596</u>	A _s G _s A _s T _s g _s a _s c _s c _s t _s c _s c _s a _s G _s A _s G _s G
			114B <u>597</u>	A _s G _s A _s T _s g _s a _s c _s c _s t _s c _s c _s a _s G _s A _s G _s g
			114C <u>598</u>	A _O G _O A _O T _O g _s a _s c _s c _s t _s c _s c _s a _s G _O A _O G _O G
			114 D <u>599</u>	a _s g _s a _s t _s g _s a _s c _s c _s t _s c _s c _s a _s g _s a _s g _s g

1585	115	TTCTCAGGAACAGCCG	115 A 600	T _s T _s C _s T _s c _s a _s g _s g _s a _s a _s c _s a _s G _s C _s C _s G
			115B 601	T _s T _s C _s T _s c _s a _s g _s g _s a _s a _s c _s a _s G _s C _s C _s g
			115C 602	T _O T _O C _O T _O c _s a _s g _s g _s a _s a _s c _s a _s G _O C _O C _O G
			115 D 603	t _s t _s c _s t _s c _s a _s g _s g _s a _s a _s c _s a _s g _s c _s c _s g
1601	116	ATGACAGGCTTTTAT	116 A 604	A _s T _s G _s A _s c _s a _s g _s g _s c _s t _s t _s T _s T _s A _s T
			116B 605	A _s T _s G _s A _s c _s a _s g _s g _s c _s t _s t _s T _s T _s A _s t
			116C 606	A _O T _O G _O A _O c _s a _s g _s g _s c _s t _s t _s T _O T _O A O _T
			116 D 607	a _s t _s g _s a _s c _s a _s g _s g _s c _s t _s t _s t _s a _s t

Please amend Table 2 starting on page 84 as follows:

Table 2 Oligomeric compounds of the invention

Oligomeric compounds were evaluated for their potential to knockdown Survivin mRNA in 15PC3 cells. The data are presented as percentage downregulation relative to mock transfected cells. Transcript steady state was monitored by Real-time PCR and normalised to the GAPDH transcript steady state. Note that all LNA C are 5'-Methyl-Cytosine.

Target site	Seq ID NO:	Oligomeric compound Sequence 5'-3'	Seq ID+ Design NO:	Specific design of Oligomeric compound Capital letters β-D-oxy-LNA s=phosphorthioate O=O-P(O) ₂ -O- Small letters DNA sugar	% Inhibition at 25 nM	% Inhibition at 5 nM oligo.
62(c)	117	AGGCAGGGGGCAACGT	117 A 608	A _s G _s G _s C _s a _s g _s g _s g _s g _s c _s a _s A _s C _s G _s T	<20	<20
			117 B	A _s G _s G _s C _s a _s g _s g _s g _s g _s c _s a _s A _s C _s G _s t		

			<u>609</u>			
			<u>117</u> € <u>610</u>	A ₀ G ₀ G ₀ C ₀ a _s g _s g _s g _s g _s c _s a _s A ₀ C ₀ G ₀ T		
			<u>117</u> D <u>611</u>	a _s g _s g _s c _s a _s g _s g _s g _s g _s c _s a _s c _s g _s t		
119(c)	118	CCAAGAAGGGCCAGTT	<u>118</u> A <u>612</u>	C _s C _s A _s A _s g _s a _s a _s g _s g _s c _s c _s A _s G _s T _s T	87	33
			<u>118</u> B <u>613</u>	C _s C _s A _s A _s g _s a _s a _s g _s g _s c _s c _s A _s G _s T _s t		
			<u>118</u> € <u>614</u>	C ₀ C ₀ A ₀ A ₀ g _s a _s a _s g _s g _s c _s c _s A ₀ G ₀ T ₀ T		
			<u>118</u> D <u>615</u>	c _s c _s a _s a _s g _s a _s a _s g _s g _s c _s c _s a _s g _s t _s t		
190(c)	119	TGGCTCGTTCTCAGTG	<u>119</u> A <u>616</u>	T _s G _s G _s C _s t _s c _s g _s t _s t _s c _s t _s c _s A _s G _s T _s G	79	27
			<u>119</u> B <u>617</u>	T _s G _s G _s C _s t _s c _s g _s t _s t _s c _s t _s c _s A _s G _s T _s g		
			<u>119</u> € <u>618</u>	T ₀ G ₀ G ₀ C ₀ t _s c _s g _s t _s t _s c _s t _s c _s A ₀ G ₀ T ₀ G ₀		
			<u>119</u> D <u>619</u>	t _s g _s g _s c _s t _s c _s g _s t _s t _s c _s t _s c _s a _s g _s t _s g		
193(c)	120	GTCTGGCTCGTTCTCA	<u>120</u> A <u>620</u>	G _s T _s C _s T _s g _s g _s c _s t _s c _s g _s t _s t _s C _s T _s C _s A	84	47
			<u>120</u> B <u>621</u>	G _s T _s C _s T _s g _s g _s c _s t _s c _s g _s t _s t _s C _s T _s C _s a		
			<u>120</u> € <u>622</u>	G ₀ T ₀ C ₀ T ₀ g _s g _s c _s t _s c _s g _s t _s t _s C ₀ T ₀ C ₀ A		
			<u>112</u> D	g _s t _s c _s t _s g _s g _s c _s t _s c _s g _s t _s t _s c _s t _s c _s a		

			<u>623</u>			
194(c)	121	AGTCTGGCTCGTTCTC	<u>121</u> A <u>624</u>	A _s G _s T _s C _s t _s g _s g _s c _s t _s c _s g _s t _s T _s C _s T _s C	75	49
			<u>121</u> B <u>625</u>	A _s G _s T _s C _s t _s g _s g _s c _s t _s c _s g _s t _s T _s C _s T _s c		
			<u>121</u> C <u>626</u>	A ₀ G ₀ T ₀ C ₀ t _s g _s g _s c _s t _s c _s g _s t _s T ₀ C ₀ T ₀ C		
			<u>121</u> D <u>627</u>	a _s g _s t _s c _s t _s g _s g _s c _s t _s c _s g _s t _s t _s c _s t _s c		
168(c)	122	TGGATGAAGCCAGCCT	<u>122</u> A <u>628</u>	T _s G _s G _s A _s t _s g _s a _s g _s c _s c _s a _s G _s C _s C _s T	67	41
			<u>122</u> B <u>629</u>	T _s G _s G _s A _s t _s g _s a _s g _s c _s c _s a _s G _s C _s C _s t		
			<u>122</u> C <u>630</u>	T ₀ G ₀ G ₀ A ₀ t _s g _s a _s g _s c _s c _s a _s G ₀ C ₀ C ₀ T		
			<u>122</u> D <u>631</u>	t _s g _s g _s a _s t _s g _s a _s g _s c _s c _s a _s g _s c _s c _s t		
215(c)	123	AGCAGAAGAAACTG	<u>123</u> A <u>632</u>	A _s G _s C _s A _s g _s a _s g _s a _s a _s c _s A _s C _s T _s G	85	26
			<u>123</u> B <u>633</u>	A _s G _s C _s A _s g _s a _s g _s a _s a _s c _s A _s C _s T _s g		
			<u>123</u> C <u>634</u>	A ₀ G ₀ C ₀ A ₀ g _s a _s g _s a _s a _s c _s A ₀ C ₀ T ₀ G		
			<u>123</u> D <u>635</u>	a _s g _s c _s a _s g _s a _s g _s a _s a _s c _s a _s c _s t _s g		
261(c)	124	TCCTCTATGGGGTCGT	<u>124</u> A <u>636</u>	T _s C _s C _s T _s c _s t _s a _s t _s g _s g _s g _s T _s C _s G _s T	23	<20
			<u>124</u> B	T _s C _s C _s T _s c _s t _s a _s t _s g _s g _s g _s T _s C _s G _s t		

			637			
			124 € 638	T ₀ C ₀ C ₀ T ₀ c _s t _s a _s t _s g _s g _s g _s T ₀ C ₀ G ₀ T		
			124 D 639	t _s c _s c _s t _s c _s t _s a _s t _s g _s g _s g _s t _s c _s g _s t		
286(c)	125	GCAACCGGACGAATGC	125 A 640	G _s C _s A _s A _s c _s c _s g _s g _s a _s c _s g _s a _s A _s T _s G _s C	64	<20
			125 B 641	G _s C _s A _s A _s c _s c _s g _s g _s a _s c _s g _s a _s A _s T _s G _s c		
			125 € 642	G ₀ C ₀ A ₀ A ₀ c _s g _s g _s a _s c _s g _s a _s A ₀ T ₀ G ₀ C		
			125 D 643	g _s c _s a _s a _s c _s c _s g _s g _s a _s c _s g _s a _s a _s t _s g _s c		
267(c)	126	TTATGTTCTCTATGG	126 A 644	T _s T _s A _s T _s g _s t _s t _s c _s c _s t _s c _s t _s A _s T _s G _s G	53	<20
			126 B 645	T _s T _s A _s T _s g _s t _s t _s c _s c _s t _s c _s t _s A _s T _s G _s g		
			126 € 646	T ₀ T ₀ A ₀ T ₀ g _s t _s t _s c _s c _s t _s c _s t _s A ₀ T ₀ G O ₀ G		
			126 D 647	t _s t _s a _s t _s g _s t _s t _s c _s c _s t _s c _s t _s a _s t _s g _s g		
325(c)	127	GGTTAATTCTTCAAAC	127 A 648	G _s G _s T _s T _s a _s a _s t _s t _s c _s t _s c _s A _s A _s A _s C	17	<20
			127 B 649	G _s G _s T _s T _s a _s a _s t _s t _s c _s t _s c _s A _s A _s A _s c		
			127 € 650	G ₀ G ₀ T ₀ T ₀ a _s a _s t _s t _s c _s t _s c _s A ₀ A ₀ A O ₀ C		
			127 D	g _s g _s t _s t _s a _s a _s t _s t _s c _s t _s c _s a _s a _s c		

			<u>651</u>			
353(c)	128	CTCTGTCCAGTTTCAA	128 A <u>652</u>	C _s T _s C _s T _s g _s t _s c _s c _s a _s g _s t _s T _s C _s A _s A	76	60
			128 B <u>653</u>	C _s T _s C _s T _s g _s t _s c _s c _s a _s g _s t _s T _s C _s A _s a	77	
			128 C <u>654</u>	C _O T _O C _O T _O g _s t _s c _s c _s a _s g _s t _s T _O C _O A O _A		
			128 D <u>655</u>	c _s t _s c _s t _s g _s t _s c _s c _s a _s g _s t _s t _s c _s a _s a		
375(c)	129	GCAATTTTGTTCTTGG	129 A <u>656</u>	G _s C _s A _s A _s t _s t _s t _s g _s t _s c _s T _s T _s G _s G	73	49
			129 B <u>657</u>	G _s C _s A _s A _s t _s t _s t _s g _s t _s c _s T _s T _s G _s g		
			129 C <u>658</u>	G _O C _O A _O A _O t _s t _s t _s g _s t _s c _s T _O T _O G O _G		
			129 D <u>659</u>	g _s c _s a _s a _s t _s t _s t _s g _s t _s c _s t _s t _s g _s g		
464(c)	130	CTCAATCCATGGCAGC	130 A <u>660</u>	C _s T _s C _s A _s a _s t _s c _s c _s a _s t _s g _s g _s C _s A _s G _s C	77	40
			130 B <u>661</u>	C _s T _s C _s A _s a _s t _s c _s c _s a _s t _s g _s g _s C _s A _s G _s c		
			130 C <u>662</u>	C _O T _O C _O A _O a _s t _s c _s c _s a _s t _s g _s g _s C _O A _O G _O C		
			130 D <u>663</u>	c _s t _s c _s a _s a _s t _s c _s c _s a _s t _s g _s g _s c _s a _s g _s c		
159(c)	131	CCAGCCTCGGCCATCC	131 A <u>664</u>	C _s C _s A _s G _s c _s c _s t _s c _s g _s g _s c _s c _s A _s T _s C _s C	80	29
			131 B	C _s C _s A _s G _s c _s c _s t _s c _s g _s g _s c _s c _s A _s T _s C _s c	94	

			<u>665</u>			
			<u>131</u> € <u>666</u>	C₀C₀A₀G₀c_st_sc_sg_sg_sc_sc_sA₀T₀ C₀C		
			<u>131</u> D <u>667</u>	c_sc_sa_sg_sc_sc_st_sc_sg_sg_sc_sa_st_sc_sc		
350(c)	132	TGTCCAGTTTCAAAAA	<u>132</u> A <u>668</u>	T_sG_sT_sC_sc_sa_sg_st_st_sc_sa_sA_sA_sA_sA	<20	<20
			<u>132</u> B <u>669</u>	T_sG_sT_sC_sc_sa_sg_st_st_sc_sa_sA_sA_sA_sa		
			<u>132</u> € <u>670</u>	T₀G₀T₀C₀c_sa_sg_st_st_sc_sa_sA₀A₀A O_A		
			<u>132</u> D <u>671</u>	t_sg_st_sc_sc_sa_sg_st_st_sc_sa_sa_sa_sa		
351(c)	133	CTGTCCAGTTTCAAAA	<u>133</u> A <u>672</u>	C_sT_sG_sT_sc_sc_sa_sg_st_st_sc_sA_sA_sA_sA	<20	<20
			<u>133</u> B <u>673</u>	C_sT_sG_sT_sc_sc_sa_sg_st_st_sc_sA_sA_sA_sa		
			<u>133</u> € <u>674</u>	C₀T₀G₀T₀c_sa_sg_st_st_sc_sA₀A₀A O_A		
			<u>133</u> D <u>675</u>	c_st_sg_st_sc_sc_sa_sg_st_st_sc_sa_sa_sa		
47(c)	134	TCGGGGCACCCATGCC	<u>134</u> A <u>676</u>	T_sC_sG_sG_sg_sg_sc_sa_sc_sc_sc_sa_sT_sG_sC_s C		
			<u>134</u> B <u>677</u>	T_sC_sG_sG_sg_sg_sc_sa_sc_sc_sc_sa_sT_sG_sC_s c		
			<u>134</u> € <u>678</u>	T₀C₀G₀G₀g_sg_sc_sa_sc_sc_sc_sa_sT₀ G₀C₀C		
			<u>134</u> D	t_sc_sg_sg_sg_sg_sc_sa_sc_sc_sc_sa_st_sg_sc_sc		

			<u>679</u>	
456(c)	135	ATGGCAGCCAGCTGCT	<u>135</u> A <u>680</u>	A _s T _s G _s G _s c _s a _s g _s c _s c _s a _s g _s c _s T _s G _s C _s T
			<u>135</u> B <u>681</u>	A _s T _s G _s G _s c _s a _s g _s c _s c _s a _s g _s c _s T _s G _s C _s t
			<u>135</u> C <u>682</u>	A ₀ T ₀ G ₀ G ₀ c _s a _s g _s c _s c _s a _s g _s c _s T ₀ G ₀ C ₀ T
			<u>135</u> D <u>683</u>	a _s t _s g _s g _s c _s a _s g _s c _s c _s a _s g _s c _s t _s g _s c _s t
470(c)	136	AGAGGCCTCAATCCAT	<u>136</u> A <u>684</u>	A _s G _s A _s G _s g _s c _s c _s t _s c _s a _s a _s t _s C _s C _s A _s T
			<u>136</u> B <u>685</u>	A _s G _s A _s G _s g _s c _s c _s t _s c _s a _s a _s t _s C _s C _s A _s t
			<u>136</u> C <u>686</u>	A ₀ G ₀ A ₀ G ₀ g _s c _s c _s t _s c _s a _s a _s t _s C ₀ C ₀ O ₀ A ₀ T
			<u>136</u> D <u>687</u>	a _s g _s a _s g _s g _s c _s c _s t _s c _s a _s a _s t _s c _s c _s a _s t
55(c)	137	GGGCAACGTCGGGGCA	<u>137</u> A <u>688</u>	G _s G _s G _s C _s a _s a _s c _s g _s t _s c _s g _s g _s G _s G _s C _s A
			<u>137</u> B <u>689</u>	G _s G _s G _s C _s a _s a _s c _s g _s t _s c _s g _s g _s G _s G _s C _s a
			<u>137</u> C <u>690</u>	G ₀ G ₀ G ₀ C ₀ a _s a _s c _s g _s t _s c _s g _s g _s G ₀ G ₀ C ₀ A
			<u>137</u> D <u>691</u>	g _s g _s g _s c _s a _s a _s c _s g _s t _s c _s g _s g _s g _s c _s a
66(c)	138	TGCCAGGCAGGGGGCA	<u>138</u> A <u>692</u>	T _s G _s C _s C _s a _s g _s g _s c _s a _s g _s g _s G _s G _s C _s A
			<u>138</u> B	

			<u>693</u>	
			<u>138</u> C <u>694</u>	T _s G _s C _s C _s a _s g _s g _s c _s a _s g _s g _s g _s G _s G _s C _s A
			<u>138</u> D <u>695</u>	t _s g _s c _s c _s a _s g _s g _s c _s a _s g _s g _s g _s g _s c _s a
140(c)	139	CCGGGGTGCAGGCGCA	<u>139</u> A <u>696</u>	C _s C _s G _s G _s g _s g _s t _s g _s c _s a _s g _s g _s C _s G _s C _s A
			<u>139</u> B <u>697</u>	C _s C _s G _s G _s g _s g _s t _s g _s c _s a _s g _s g _s C _s G _s C _s a
			<u>139</u> C <u>698</u>	C _O C _O G _O G _O g _s g _s t _s g _s c _s a _s g _s g _s C _O G _O C _O A
			<u>139</u> D <u>699</u>	c _s c _s g _s g _s g _s g _s t _s g _s c _s a _s g _s g _s c _s g _s c _s a
148(c)	140	CATCCGCTCCGGGGTG	<u>140</u> A <u>700</u>	C _s A _s T _s C _s c _s g _s c _s t _s c _s c _s g _s g _s G _s G _s T _s G
			<u>140</u> B <u>701</u>	C _s A _s T _s C _s c _s g _s c _s t _s c _s c _s g _s g _s G _s G _s T _s g
			<u>140</u> C <u>702</u>	C _O A _O T _O C _O c _s g _s c _s t _s c _s c _s g _s g _s G _O G _O T _O G
			<u>140</u> D <u>703</u>	C _s A _s T _s C _s C _s G _s C _s T _s C _s C _s G _s G _s G _s G _s sT _s G
177(c)	141	GTGGGGCAGTGGATGA	<u>141</u> A <u>704</u>	G _s T _s G _s G _s g _s g _s c _s a _s g _s t _s g _s g _s A _s T _s G _s A
			<u>141</u> B <u>705</u>	G _s T _s G _s G _s g _s g _s c _s a _s g _s t _s g _s g _s A _s T _s G _s a
			<u>141</u> C <u>706</u>	G _O T _O G _O G _O g _s g _s c _s a _s g _s t _s g _s g _s A _O T _O G _O A
			<u>141</u> D	g _s t _s g _s g _s g _s g _s c _s a _s g _s t _s g _s g _s a _s t _s g _s a

			<u>707</u>	
260(c)	142	CCTCTATGGGGTCGTC	142 A <u>708</u>	C _s C _s T _s C _s t _s a _s t _s g _s g _s g _s g _s t _s C _s G _s T _s C
			142 B <u>709</u>	C _s C _s T _s C _s t _s a _s t _s g _s g _s g _s g _s t _s C _s G _s T _s t
			142 C <u>710</u>	C ₀ C ₀ T ₀ C ₀ t _s a _s t _s g _s g _s g _s g _s t _s C ₀ G O ₀ T ₀ C
			142 D <u>711</u>	c _s c _s t _s c _s t _s a _s t _s g _s g _s g _s g _s t _s c _s g _s t _s c
274(c)	143	ATGCTTTTATGTTCC	143 A <u>712</u>	A _s T _s G _s C _s t _s t _s t _s t _s a _s t _s g _s T _s T _s C _s C
			143 B <u>713</u>	A _s T _s G _s C _s t _s t _s t _s t _s a _s t _s g _s T _s T _s C _s t
			143 C <u>714</u>	A ₀ T ₀ G ₀ C ₀ t _s t _s t _s t _s a _s t _s g _s T ₀ T ₀ C ₀ C
			143 D <u>715</u>	a _s t _s g _s c _s t _s t _s t _s t _s a _s t _s g _s t _s c _s c
384(c)	144	GTTTCCTTTGCAATTT	144 A <u>716</u>	G _s T _s T _s T _s c _s c _s t _s t _s g _s c _s a _s A _s T _s T _s T
			144 B <u>717</u>	G _s T _s T _s T _s c _s c _s t _s t _s g _s c _s a _s A _s T _s T _s t
			144 C <u>718</u>	G ₀ T ₀ T ₀ T ₀ c _s c _s t _s t _s g _s c _s a _s A ₀ T ₀ O ₀ T ₀ T
			144 D <u>719</u>	g _s t _s t _s c _s c _s t _s t _s g _s c _s a _s a _s t _s t
ISIS 23722	145	TGTGCTATTCTGTGAA TT (18-mer)	145 A <u>720</u>	T _s G _s T _s G _s c _s a _s t _s c _s t _s g _s g _s A _s A _s T _s T
			145 C <u>721</u>	T ₀ G ₀ T ₀ G _s c _s a _s t _s c _s t _s g _s g _s A ₀ A ₀ T ₀ T

			<u>721</u>	T _O T
			145 D <u>722</u>	t _s g _s t _s g _s c _s t _s a _s t _s t _s c _s t _s g _s t _s g _s a _s a _s t _s t
			145F <u>723</u>	<i>T_sG_sT_sG_sc_st_sa_st_st_sc_st_sg_st_sg_sA_sA_sT_sT</i>
	146		146 A <u>724</u>	T _s <u>A_sA_s</u> G _s c _s t _s g _s t _s t _s c _s t _s a _s t _s g _s <u>T_sG_s</u> T _s T*
			146 € <u>725</u>	T _O <u>A_OA_O</u> G _s c _s t _s g _s t _s t _s c _s t _s a _s t _s g _s <u>T_OG_O</u> T _O T*
			146F <u>726</u>	<i>T_s<u>A_sA_s</u>G_sc_st_sg_st_st_sc_st_sa_st_sg_s<u>T_sG_s</u>T_sT*</i>

* relates to compound Underlined indicates mismatch compared to above compound.
 Compound 145F and 146F contains the MOE chemistry in capital letters italic which is the compound ISIS23722.

Please amend Table 3 starting on page 88 as follows:

Table 3 IC₅₀ (nM) of LNA (β -D-oxy-LNA) containing oligomeric in two cell lines of different origin

Oligomeric compounds were evaluated for their potential to knockdown Survivin mRNA in 15PC3 and MCF7 cells. Transcript steady state was monitored by Real-time PCR and normalised to the GAPDH transcript steady state.

Seq ID/design NO	MCF7	15PC3
2A-147	28	5
2B-148		<5
4A-155		<5
4B-156		5
6A-163	8	3
6B-164		<5
9A-175	11	3
15A-199	1	<1
15B-200		<1
15E-203		1
118A-612		<5
120A-620		<25
123A-623		<5
128A-652		<5
128B-653		<25
129A-656		<25
131A-664		<25
131B-665		<5

Please amend the paragraph on page 88, lines 14-15, as follows:

Compounds of particular interest are ~~2A, 2B, 4A, 4B, 6A, 6B, 15A, 15B, 15E, 119A, 119B, 121A, 121B, 128A, 128B, 130A, 130B, 131A and 131B.~~ 147, 148, 155, 156, 163, 164, 199, 200, 203, 616, 617, 624, 625, 652, 653, 660, 661, 664, and 665.

Please amend the paragraph on page 95, lines 5-21, as follows:

Human 15PC3 xenografted tumors according to Example 13 were homogenized in 10 volumes of 0,5% Igepal CA-630, 25 mM Tris pH 8.0, 25 mM EDTA, 100 mM NaCl, 1mg/ml Proteinase K1 and incubated overnight at 37 degrees celsius followed by phenol-chloroform extraction. The concentration of antisense oligonucleotide 2650 in the combined aqueous phase was determined using a sequence specific ELISA assay. Two probes, one labelled with biotin and one labelled with digoxigenin (DIG) with complementary sequences to the antisense oligonucleotide are hybridised to the antisense oligo. The complex is captured by immobilized streptavidin and quantified using a horse raddish peroxidase-conjugated anti-digoxigenin antibody and standard ELISA procedures. Briefly, 10 nM DNA capture probe (5'-aactgtgc-Biotin-3') and 10 nM LNA detection probe (5'-DIG-GATGTTTCgatgttcc-3')(SEQ ID NO: 738) were mixed with sample or standards in 1 % blocking reagent (Roche cat. 1 096 176) in PBS. The probes were annealed to the oligo by heating the mixture to 70 degrees celsius and gradual cooling to 20 degrees Celsius. The mixture was transferred to streptavidin-coated wells. The amount of captured DIG-probe is quantified using an HRP-conjugated Anti-DIG antibody fragment (Roche) and standard ELISA procedures. At least 1,3µg/g tumours tissue of the oligomeric compound 15A was detected (data not adjusted for recovery).